	4	140117			
			NM 052885		16 CC0201
	64224 EI 122				T
	51095 TRNT1				
	64326 RFWD2	NP 071902 6			
	135398 C6orf141	NP_699175 1	NM_153344	40	1
	58484 CARD12	NP_067032 5	NM_021209		
	142680 SLC34A3	NF 343133			
	80144 FRASI			45	
	7579 ZNF31				1557502 or
	246269 LACE1	NP 660358 2			
		NP_036419 2			at
248	126248 LOC	NP_775750			
	317719 KLHL10	NP_689680 3			1552425 a at
19	254173 FLJ36119	NP 694986			1552425 at
	5150 PDE7A	NP_002595			1552343 s at
	145258 GSC	NP_776248			18 0CC2CC1
AP1 GTPase, IMAP family member 1	75		NM_130759		1552316_a_at
		7 816500 dN	NM_005927	4D	1552312 a at
29937	7	NP_653198	NM_144597	4D	1552310 at
	65		NM_017861	4D	1552291 at
	172 AFG3L1	NP_001123	NM_001132	4D	1552287_s_at
ATP6V1E2 ATPase, H+ transporting, lysosomal 31kDa, V1	90423 ATP	NP_542384	NWI_080603	ŧ	. 00 mm 00 mt
MGC17337 similar to RIKEN cDNA 5730528L13 sene	91283 MGG				18 B / /7701
	54899 PXK	NP_060241	NM 017771		18 +177CC1
	5594 MAPK1		/C69C1 TAIN		W The Courses
	5594 MAPKI		NIM 120057		1552264 a at
DC2 WAP fourdisulfide core domain 2	10406 WFDC2		NIM 120057		1552263 at
77			NM 080735		1552261 at
N21 protein tyrosine phosphatase, nonreceptor type 21	11099 PTPN21	NF 0089/0	MM 057671		1552258 at
paired box gene 8			NM 007039	40	1320 at
Gene Symbol	LocusLink Gen		Experimental HumanProtAcces	cyberment	17 of the lot

RAS and EF hand domain containing	RASEF	8C18C1	INF 069/60	102010	i	1
RAS and EF hand domain containing	KASEF	120120	NID 600700	NM 152573	4D	24
phosphodiesterase 5A, cGMPspecific	FUESA	150150	ND 690796	NM 152573	4D	
inspenience protein MOC14126	DDES A	7598	NP 737773	NM 033431	4D	1553175 s at
hypothetical passes MCC1 1132	MGC14126	84984	NP 116287	NM_032898	4D	1553158_at
ATPase, H+ transporting, lysosomal 38kDa, V0 subun	ATP6V0D2	245972	NF 0897/8	NM 132303	100	1
hypothetical protein FLJ25422	FLJ25422	202151	NP 659437	142668 MIN	45	
chromosome 9 open reading frame 72	C9orf72	203228	NP 659442	NIN 145000	100	
RBassociated KRAB repressor	RBAK	57786	NP 066986	COLIZO TAIN	4D	
nuclear transcription factor, Xbox binding	NFX1	4799	NP 66/345	14/162 14/14/	40	1553122 at
tigger transposable element derived 1	TIGDI	200765	NP_663748	NM 145/02	45	1553103 at
BCL2like 11 (apoptosis facilitator)	BCL2L11	10018	NP_619533	NM_138627	ŧ	18 S 0000 C
BCL2like 11 (apoptosis facilitator)	BCL2L11	10018	NP_619532	NM_138626	45	1553006 a at
				NM_032362	4D	1553087_at
ankvrin reneat and SOCS becomes in 10	ASB10	136371	NP_543147	NM_080871	4D	1553039 a at
Synantotagominilibe 5	SYTL5	94122	NP_620135	NM_138780	40	1553033_at
sulfite oxidase	SUOX	6821	NP_000447	NM_000456	4D	1553030 a at
aminosoming oncogonence i				NM_054029	4D	1552955_at
mel transforming oncognostic 1	MELLI	79258	NP_258428	NM_033467	4D	1552930 at
fidoetiniika i	FIGNL	63979	NP_071399	NM_022116	4D	1552921_a_at
interlenkin 23 recentor	IL23R	149233	NP_653302	NM_144701	4D	1552912 a at
chromosome 10 open reading frame 27	C10orf27	219793	NP 689923	NM_152710	4D	155288/ at
atonal homolog 7 (Droscobile)	АТОН7	220202	NP_660161	NM_145178	4D	1552879 a at
Shydroxytryntamina (sarotonia)	HTR6	3362	NP_000862	NM_000871	4D	1552857 a at
Zinc finger protein 610	ZNF619	285267	NP_775927	NM_173656	4D	1552836_at
tolllike recentor 4	TLR4	7099	NP_612567	NM_138557	4b	1552798 a at
and (series) D				NM_153039	4D	1552789_at
helicase (DNA) B	HELB	92797	NP_387467	NM 033647	4D	1552788 a at
Ctype lectin domain family 1	CLEC4D	338339	NP_525126	NM_080387	4D	1552772_at
calcium/calmodulindependent protein kinase	CAMKKI	84254	NP_757344	NM_1/2207	ŧ	1002100 at
MCF.2 cell line derived transforming segmencelike	MCF2L	23263	NP 079255	NM 024979	ŧ	1557760 ct
WW domain containing E3 ubiquitin protein ligase 2	WWP2	11060	NP_008945	NM 007014	ŧ	1002/07 S at
					;	1000000

1553193 at	40	NM_152355	NP_689568	126068	ZNF441	zinc finger protein 441
1000218 a at	ŧ	NM 032434	NP 115810	84450	ZNF512	zinc finger protein 512
1552260 -t	ŧ	NM 152633	NP 689846	2187	FANCB	Fanconi anemia, complementation group B
1553269 at	4	NM_153691	NP_710158	255403	FLJ90036	hypothetical protein FLJ90036
1553272_at	‡ D	NM_078483	NP_510968	206358	SLC36A1	solute carrier family 36 (proton/amino acid
1553296_at	4D	NM_032787	NP_116176	84873	GPR128	G proteincoupled recentor 128
553297_a_at	4D	NM_172313	NP_758519	1441	CSF3R	colony stimulating factor 3 recentor (granulogic
553311_at	45	NM_173644	NP_775915	284756	FLJ33860	hypothetical protein FLJ33860
1553328_a_at	4D	NM_003054	NP_003045	6571	SLC18A2	solute carrier family 18 (vesicular monoamine),
1553363_at	4D	NM_152554	NP 689767	154386	C60rf195	chromosome 6 open wording from
553373_at	4D	NM_144625	NP 653226	128025	FI J32978	hypothetical protein El 122026
553400 a at	4D	NM_152466	NP 689679	147081	FLJ25168	hynothetical protein El 125169
553422 s at	4D	NM_145892	NP_665899	54715	A2BP1	ataxin 2binding protein 1
1333423 a at	40	NM_144682	NP_653283	146857	FLJ31952	hypothetical protein FLJ31952
18 octoo	ŧ	NM_1/35/8	NP_775849	283135	FLJ90834	hypothetical protein FLJ90834
553512 a at	ŧ	NM 053016	NP_443749	114299	PALM2	paralemmin 2
553576 at	ŧ	NM 17860	NP 776272	3228	HOXC12	homeo box C12
02000 00	ŧ	NM 1/0011	NP_/89/81	126205	NALP8	NACHT, leucine rich repeat and PYD containing
1553539 at	4D	NM_175053	NP_778223	121391	K6IRS4	keratin 6 irs4
553550 at	4D	NM_173858	NP_776257	317705	VNIRS	vomeronasai i recentor s
1553568 a at	45	NM_005323	NP 005314	3010	HISTIHIT	histone 1, H1t
1553574 at	4D	NM_176891	NP_795372	338376	IFNEI	interferon epsilon i
553579_a_at	4b	NM_058206	NP_478113	10407	SPAGII	sperm associated antigen []
1003626 a at	40	NM_152347	NP_689560	124989	FLJ40342	hypothetical protein FLJ40342
1553677 a at	4D	NM_152902	NP_690866	261726	MGC3794	putative MAPK activating protein
553685 s at	4D	NM_138473	NP 612482	6667	SPI	Spl transcription factor
553703_at	4D	NM_153358	NP_699189	163049	FLJ90396	hypothetical protein FL 190396
553/04 x at	40	NM_153358	NP 699189	163049	FLJ90396	hypothetical protein FLJ90396
333723 at	40	NM_170776	NP_740746	222487	GPR97	G proteincoupled receptor 97
555/68 a at	40	NM_173674	NP_775945	285761	DCBLDI	discoidin, CUB and LCCL domain containing 1
3337/0 a at	÷	NM 033438	NP 254273	89886	SLAMF9	SI AM familia month

neuroligin 2 WINS1 protein with Drosophila Lines (Lin)	Transi pro						
2	WING pro	WINSI	55180	NP_859523	WM_181739	÷	1004400_at
	neuroligin 2	NLGN2	57555	NP 065846	NM 020795	1 5	55/1/55 at
elegans	elegans			1		ð	001155
opposition plotein 11.320123	antarior pho	APHIA	51107	NP 057106	NM_016022	4D	554417 s at
protein El 12012s	hypothetica	FLJ20125	54826	NP_060146	NM_017676	4D	1554345 a at
ase HEL308	DNA helicase HEL 308	HEL308	113510	NP_598375	NM_133636	4D	1554342 s at
cetylase 4	histone deacetylase 4	HDAC4	9759	NP_006028	NM_006037	40	1334322 a at
hypothetical protein LOC136306	hypothetica	LOC136306	136306	NP_777619	NM_174959	÷	1554300 a at
protein	adult retina protein	LOC153222	153222	NP_705835	NM 153607	ŧ	1554229 at
PHD finger protein 21A	PHD finger	PHF21A	51317	NP_057705	NM_016621	40	1004103 a at
unc51like kinase 2 (C. elegans)	unc51like k	ULK2	9706	NP_055498	NM_014683	4	1554112 a at
amyotrophic lateral sclerosis 2 (invenile) chromos	amyotrophi	ALS2CR16	130029	NP_991106	NM_205543	4b	1554108 at
DEAD (AspGluAlaAsp) box polymentide \$1	DEAD (Asi	DDX51	317781	NP_778236	NM_175066	4D	1554085 at
serine palmitovltransferase, long chain base subus	serine palm	SPTLCI	10558	NP 847894	NM_178324	45	1004003 at
zinc finger protein 12 (KOX 3)	zinc finger	ZNF12	7559	NP_008887	NM_006956	4	1554021 a at
apoptosisrelated protein PNASI	apoptosisre	FLJ39616	51275	NP_057618	NM_016534	45	155598/ at
chromosome 20 open reading frame 161	chromoson	C20orf161	90203	NP_690857	NM_152897	ŧ	155300 at
UDPGal:betaGal beta 1,3galactosyltransferase polyp	UDPGat:be	B3GALT6	126/92	NF_342172	N. 15000	1	\$\$2060 at
omponent 6	exosome component 6	EXOSC6	118400	NE 4/61/20	NIM 090005	4	1553959 a at
			+	NID 470106	01C850 MIN	4D	1553947 at
hypothetical protein FLJ40365	nypometica	1 L340303	10000		NM 131915	4D	1553900 s at
b proteincoupled receptor NRGX2	o proteino	El Moses	126402	NP 775753	NM 173482	₽D	1553893_at
Description of the property of	G proteins	MRGX2	117194	NP 473371	NM_054030	4D	1553889_at
hynothetical protein MCC35220	hypothetic	MGC35338	130619	NP 689602	NM_152389	4D	1553886_at
al protein MGC24712	hypothetic	MGC34713	285600	NP_775936	NM_173665	4D	1553868 a at
emonitorito open reachig mane 64	01100001				NM_138339	4D	1553865 a at
chromosome 10 open reading frame 64	chromoson	C10orf64	159491	NP_775795	NM 173524	₽ ₽	1553863_at
al protein MGC40368	hypothetic	MGC40368	255394	NP_689985	NM_152772	4D	1553861 at
Cohen syndrome 1	Cohen syndrome	COHI	157680	NP_689777	NM_152564	4D	1553852_at
ne 21 onen reading frame 00	chromosor	C21orf88	114041	NP_715635	NM_153754	4D	1553850 at
hypothetical protein El 125371	hypothetic	FLJ25371	152940	NP 689756	NM 152543	40	1553838 at
SRY (sex determining region Valor 2	SRY (sex	SOX3	6658	NP_005625	NM_005634	4D	1553802 a at
					NM_032202	4D	1553793 a at

Nacylsphingosine amidohydrolase (acid ceramidase) ceramidase) transmembrane proteine de			TO COMMO	NIM 000770	4D	1555497 a at
Nacylsphingosine amidohydrolase (acid ceramidase) transmembrane protein 1 hypothetical gene supported by BC046200	SDCCAG10	10283	NP_005860	NM 005869	40	
Nacylsphingosine amidohydrotase (acid ceramidase) transmembrane protein 1	LOC388407	388407	NP 982249	NM 203425	ŧ	L
Nacylsphingosine amidohydrolase (acid ceramidase)	TMEMI	7109	NP_001001723	NM 001001723	5 6	s at
The same of the same of	ASAHI	427	NP_004306	NM_004515	ŧ	2
FGDI family, member 3	FGD3	6	NP_149077	NM 033086	40	
ATPbinding cassette, subfamily B (MDR/TAP), member	ABCB5	340273	NP_848654	NW_1/8339	ŧ	
KIAA0377 gene product	KIAA0377	9677	NP 055474	NM 014659	ŧ	ä
retinitis pigmentosa Hike	RPILI	94137	NP 849188	NM 178857	ŧ	
hypothetical gene supported by BC055092	LOC401466	401466	XP_376783	XM_376783	#	
hypothetical gene supported by BC055092	LOC401466	401466	XP_376783	XM_376783	4D	
four and a half LIM domains 5	FHL5	9457	NP_065228	NM_020482	4D	at
preBcell colony enhancing factor 1	PBEFI	10135	NP 877591	NM_182/90	#	
quaking homolog, KH domain RNA binding (mouse)	QKI	9444	NP_006766	NWI_006//5	ŧ	1
hypothetical protein MGC40069	MGC40069	348035	NP_872421	NM 182615	ŧ	
scavenger receptor class F, member 1	SCARFI	8578	NP_663324	NM_145349	4	555073 a at
zinc finger protein 398	ZNF398	57541	NP_733787	NM 170686	ŧ	55502 a at
hypothetical protein LOC403312	MGC39545	403312	NP 982277	NM_203452	ŧ	555015 at
phospholipase A2, group IVD (cytosolic)	PLA2G4D	283748	NP 828848	NM_178034	ŧ	555002 at
RAB39, member RAS oncogene family	RAB39	54734	NP 059986	NM 017516	ŧ	554014 at
hypothetical protein FLJ32130	FLJ32130	146540	NP 689671	NM_152458	÷	1554900 X at
hypothetical protein FLJ32130	FLJ32130	146540	NP_689671	NM_152458	4	1554/69_at
transforming, acidic coiledcoil containing protein	TACC1	6867	NP_006274	NM_006283	40	1554690 a at
hypothetical protein LOC126295	LOC126295	126295	NP_775751	NM_173480	4D	1554628 at
polypyrimidine tract binding protein 2	PTBP2	58155	NP_067013	NM_021190	40	1554614 a at
cyclin M2	CNNM2	54805	NP_060119	NM_017649	ŧ	1004023 a at
growth hormone inducible transmembrane protein	GHITM	27069	NP_055209	NM_014394	ŧ	1554500 S at
regulator of Gprotein signalling 7	RGS7	6000	NP 002915	NM 002924	ŧ	1554500 a at
transmembrane protein 37	TMEM37	140738	NP_899063	NM 183240	ŧ	1554485 at
PHD finger protein 20like 1	PHF20L1	51105	NP 079154	NM 024878	÷	1004472 a at
WINS1 protein with Drosophila Lines (Lin) homologo	WINSI	55180	NP_859524	NM_181740	4	2

zinc finger protein 230 Zinc finger protein 230 In profilerical gene supported by AK093266 phosphatidylinositol3phosphatic phosphatidylinositol3phosphatic phosphatidylinositol3phosphatic phosphatidylinositol3phosphatic photein (calculation) Citation (calculation) Lipotherical protein FL12/1908 male sterifity domain containing 2 SFRS protein kinase 2 Jint Stay protein kinase 2 SFRS protein kinase 2 SFRS protein kinase 2 Jint Stay protei	NEK8 ZNF230 ZNF230 LCC400238 PIPSK3 CCHR3SYT CTNND1 FLJ21908 MLSTD2 SRPK2 PIGM ZDHHC18 NFKBIL2	400238 200576 83850 83850 1500 79657 84188 6733 284098 84243	NP_8/2633 NP_848612 NP_115659 NP_038460	NM_032283 NM_013432	40 40	
zine finger protein 230 zine finger protein 230 kppothetical gene supported by AK093266 phosphatidylinositol3phosphate/phosphatidylino chr3 synapotagmin caterin (cadherinassociated protein), delta 1 kppothetical protein FL21908 male sterility domain containing 2 SFRS protein kinase 2 phosphatidylinositol glycan, class W zine finger, DHHC domain containing 18	NER8 ZNF230 LIOC400238 PIPSK3 PIPSK3 CHR3SYT CTNND1 FLJ21908 MLSTD2 SRPK2 PIGW ZDHHC18	400238 200576 83850 83850 1500 1500 79657 79657 84188 6733 284098 84243		NM_032283	4D 8	1558327_at
zine finger protein 230 zine finger protein 230 lix pothetical gene supported by AK093266 phosphatidy linositol 3 phosphately linositol 3 phosphately linositol 4 phosphately linositol 4 phosphately linositol 2 phosphately linositol 2 protein), delta 1 lix pothetical protein FL27098 lix protein fil 2 phosphately linositol glycan, class W phosphately linositol glycan, class W	NER8 ZNF230 LOC400238 PIPSK3 CHR3SYT CTNND1 FLJ21908 MLSTD2 SRPK2 PIGW	400238 200576 200576 83850 1500 1500 1500 1500 84188 6733 284098			t	120000
zine finger protein 230 zine finger protein 230 kppotheteal gene supported by AK093266 phosphatidy linositol 2phosphatie phosphatidy lino phosphatidy linositol 2phosphatie phosphatidy lino clara Synaptotagnia catetain (cadherinassociated protein), delta hypothetical protein FLI21908 male sterifly domain containing 2 SFRS protein kinase 2 SFRS protein kinase 2	NEK8 ZNF230 LOC400238 PIP5K3 PIP5K3 CHR3SYT CTNND1 CTNND1 FLJ21908 MLSTD2 SRPK2	400238 200576 200576 83850 1500 1500 79657 84188 6733	NP 872633	NM 178517		1558292 s at
zinc Inger protein 230 zinc Inger protein 230 lixpothetical gene supported by AK093266 phosphatdylinosito/3phosphate/phosphatdylino ohr3 synaptotagmin caterin (cadherinassociated protein), delta 1 bypothetical protein FLJ21908 male sterilly domain containing 2	NEK8 ZNF230 LOC400238 PIPSK3 CCHR3SYT CCTNND1 FLJ21908 MLSTD2	400238 200576 200576 83850 1500 79657		NM_182691	40	1558254 s_at
zine finger protein 230 zine finger protein 230 kippolhetical gene supported by AK093266 phosphatidy linositol 3 phosphatidy lino ohr 3 ynapotagenin caterin (cadherinassociated protein), delta 1 by pothetical protein FLJ21048	NEK8 ZNF230 LOC400238 PIPSK3 CCHR3SYT CCTNND1 FLJ21908	400238 200576 200576 83850 1500 79657	NP_115604	NM_032228	40	1558014 s at
zine finger protein 230 zine finger protein 230 lis pathettal gene supported by AK093266 phosphatidy linositol 3phosphate phosphatidy lino phosphatidy linositol 3phosphate phosphatidy lino did 3 ynaptotagmin catetin (cadherinassociated protein), delfa 1	NEK8 ZNF230 LOC400238 PIP5K3 CCHR3SYT CCTNND1	400238 200576 20850 83850	NP 078880	NM_024604	40	155/984 s_at
zine finger protein 230 hypothetical gene supported by AK093266 phosphatidylinositol3phosphate/phosphatidyline o by 3ynaptotagmin	NEK8 ZNF230 LOC400238 PIP5K3	400238 200576 83850	NP 001322	NM_001331	40	155/944 s at
zinc finger protein 230 hypothetical gene supported by AK093266 phosphatidylinositol3phosphate/phosphatidylino	NEK8 ZNF230 LOC400238 PIP5K3	400238 200576	NP_114119	NM_031913	4	1557/95 at
zinc finger protein 230 hypothetical gene supported by AK093266	NEK8 ZNF230 LOC400238	400238	NP_001002881	NW 001002881	ŧ	1007/17_dt
zinc finger protein 230	NEK8 ZNF230		XP_378473	XM_378473	ŧ	133/03/ a at
Committee of the second of the	NEK8	7773	NP_006291	NM_006300	6	1557/522 at
NIMA (never in mitosis gene a) related kinase 8		284086	NP_835464	NM_178170	4	1557172_x_at
kelchlike 18 (Drosophila)	KI HI 18	23276	NP_079286	NM_025010	4D	1557165 s_at
hypothetical protein FLJ22374	FLJ22374	84182	NP_115598	NM_032222	4D	1557141_at
hypothetical protein FLJ36070	FLJ36070	284358	NP_872380	NM_182574	4D	1557091_at
hypothetical protein LOC133491	LOC133491	133491	XP_376444	XM_376444	45	155/056 at
Lmy						
similar to Zinc finger protein RIF(Reservance)	LOC440971	440971	XP_496651	XM_496651	4D	1556744_a_at
hypothetical protein LOC283687	LOC283687	283687	NP_787094	NM_175898	4D	1556588 at
hypothetical protein I OC 197322	LOC197322	197322	NP_777577	NM_174917	4D	1556552_a_at
FGFRI oncogene partner 2	FGFR1OP2	26127	NP_056448	NM_015633	4D	1556283 s at
chromosome 10 open reading frame 90	C10orf90	118611	NP_001004298	NM_001004298	4D	1556200 a at
peroxisome biogenesis factor 13	PEX13	5194	NP_002609	NM_002618	4	1556009_at
histidine triad nucleotide binding protein 1	HINTI	3094	NP_005331	NM_005340	4	1555961 a at
cell division cycle 25A	CDC25A	993	NP_001780	NM_001789	4D	1555772_a_at
angiotensin II receptorassociated protein	AGTRAP	57085	NP_065083	NM_020350	40	1555736 a at
CD209 antigen	CD209	30835	NP_066978	NM_021155	ť	1555729 a at
regulator of Gprotein signalling 5	RGS5	8490	NP_003608	NM 003617	1	1555/25 a at
ST3 betagalactoside alpha2,3sialvltransferase 3	ST3GAL3	6487	NP 777624	NM 174964	4D	1555702 a at
SLC2A4 regulator	SLC2A4RG	56731	NP_064446	NM_020062	4D	1555500 s at
interleukin 28 receptor, alpha (interferon, lambda	IL28RA	163702	NP 775088	NM_173065	4	1555499 a at
polypeptid					;	1555 100

myoneurin	NYNN	55892	NP 061127	NM_018657	4D	1566109 at
dual oxidase 1	DUOX1	53905	NP 059130	NM 017434	40	
hypothetical protein LOC164045	LOC164045	164045	XP_371267	XM_371267	100	
chromosome 5 open reading frame 16	C5orf16	285613	NP_776189	NM_1/3828	5 6	a
hypothetical protein FLJ32110	FLJ32110	219578	NP_857597	NM 181646	15	1564021 at
plexin D1	PLXNDI	23129	NP_055918	NM 015103	ŧ	18 / COCOC1
parvin, alpha	PARVA	55742	NP_060692	NM_018222	15	15/3/57 at
hypothetical gene supported by BC040875	LOC400456	400456	XP_378545	XM_378545	ŧ	15/2150 11067901 at
sterile alpha motif domain containing 12	SAMD12	401474	NP_997389	NM_207506	ŧ	156263/_at
hypothetical gene supported by AK097381; BC040866	LOC400680	400680	XP_378783	XM_378783	£	1502548_at
CUB and Sushi multiple domains 1	CSMD1	64478	NP_150094	NM_033225	8	1562329 at
pyruvate dehydrogenase kinase, isoenzyme 4		5166	NP_002603	NM_002612	40	1562321 at
chromosome 3 open reading frame 15		89876	NP_203528	NM_033364	4D	1561429 a at
hypothetical gene supported by BC042100	LOC440602	440602	XP_498751	XM_498751	40	1561555 at
stereocilin		161497	NP_714544	NM_153700	40	1561306 s_at
chromosome 21 open reading frame 106		23181	NP_996773	NM_206890	4D	1561286 a at
hypothetical protein LOC338579	LOC338579	338579	XP_495807	XM_495807	40	1361225_at
nitric oxide synthase I (neuronal)	NOSI	4842	NP 000611	NM_000620	40	15609 /4 s at
hypothetical protein LOC285972	LOC285972	285972	XP_379530	XM_379530	4D	1560/62 at
Rhoguanine nucleotide exchange factor		64283	XP_371755	XM_371755	40	1560348_at
laminin, alpha 3		3909	NP_937762	NM_198129	4b	1560078 at
Rho guanine nucleotide exchange factor (CEE) 12	12	23365	NP_056128	NM_015313	4D	1559833_at
hypothetical protein F1 135681		283897	NP_787096	NM_175900	4D	1559584 a at
KIAA1467 protein	467	1	XP_370682	XM_370682	4D	1559302_at
spergen		~	NP_777587	NM 174927	45	1559138_a_at
KIAA 1447 protein	447	57597	XP_371084	XM_371084	4D	1559075 s at
HLA complex group 27	HCG27	253018	NP_859068	NM 181717	4D	1559050 at
KIAA1240 protein	KIAA1240	54454	XP_039676	XM_039676	4D	1558807 at
zinc finger protein 548	ZNF548	147694	NP_690873	NM 152909	Đ	1558622 a at
zinc finger protein 621	ZNF621	285268	NP_940886	NM_198484	4D	1558620 at
LATISTM protein	LATI3TM	81893	NP 112488	NM_031211	4D	1558534 at
hypothetical protein OC112220	LOC113230	113230	XP_053966	XM_053966	4D	1558412 at
hypothetical gene supported by NM 104204	LOC439911	439911	XP_498467	XM_498467	4D	1558345 a at
						The same of the sa

1566289 at	4D	NM_017504	NP_059974	26245	OR2M4	olfactory receptor, family 2, subfamily M. member
1567107 s at	8	NM_020732	NP 065783	57492	ARIDIB	AT rich interactive domain 1B (SWIIIike)
1567238 at	ŧ ŧ	NM 003290	NP 003281	7171	TPM4	tropomyosin 4
1567284 at	40 6	NM 001004060	NP 001004686	26246	OR2L2	olfactory receptor, family 2, subfamily L, member
1568590 at	4D	NM 004311	NP 001007	20219	ABI 3	olfactory receptor, family 1, subfamily J, member
1568592 at	4D	XM 378511	XP 378511	100369	ANLS	ADPribosylation factorlike 3
1568609 s at	ŧ	NM 207400	NP 997283	388685	F1 J39739	nypothetical gene supported by BC031266
1568629_s_at	4D	NM_005027	NP_005018	5296	PIK3R2	phosphoinositide3kinase, regulatory subunit 2 (p85)
1300036_at	ŧ	XM_291016	XP_291016	339804	LOC339804	hypothetical gene supported by AK075484; BC014578
1568667 s at	40	XM_496282	XP_496282	440503	LOC440503	similar to RIKEN DNA 23100761 00
1568678 s at	4D	NM_007045	NP_008976	11116	FGFR1OP	FGFRI oncogene partner
1308/13_a_at	4	NM_015173	NP_055988	23216	TBCIDI	TBCI (tre2/USP6, BUB2, cdc16) domain family,
1568720_at	40	NM_031218	NP_112495	81931	ZNF505	zinc finger protein 505
208826 at	40	NM_031862	NP_114068	4077	NBR1	neighbor of BRC A1 gene 1
568924 a at	4D	NM_178827	NP_849149	154865	FLJ35834	hypothetical protein FL [35834
1368934 s at	#	NM_014117	NP 054836	29035	PRO0149	PRO0149 protein
15 S 20060C S at	£	NM_178229	NP_839943	128239	IQGAP3	IQ motif containing GTPase activating protein 3
15690/3_x_at	#	NM_003072	NP_003063	6597	SMARCA4	SWI/SNF related, matrix associated, actin dependen
1369102_at	45	NM_000350	NP_000341	24	ABCA4	ATPbinding cassette, subfamily A (ABC1), member 4
1569189 at	4D	NM_173810	NP_776171	283237	MGC29649	hypothetical protein MCC20640
1569206_at	4D	NM_152772	NP_689985	255394	MGC40368	hynothetical protein MCC40248
1569207 s at	4D	NM 152772	NP_689985	255394	MGC40368	hypothetical protein MGC40368
1569240 at	4D	NM_031218	NP_112495	81931	ZNF505	zinc finger protein 505
369302 at	4υ	XM_374922	XP_374922	85459	KIAA1731	KIAA1731 protein
569323 at	4D	NM_002841	NP_002832	5793	PTPRG	protein tyrosine phosphatase recentor tops C
569349 at	4D	NM_020193	NP_064578	56946	Ö	chromosome II open reading frame 30
1569408 at	4D	NM_017629	NP_060099	192670		eukarvotic translation initiation factor 20 4
	#	NM_022763	NP_073600	64778	FNDC3B	fibronectin type III domain containing 3B
1204071 at	40	NM_022787	NP 073624	64802	MNATI	nicotinomide nucleotide ed

		200	05000			1000004
RAD21 homolog (S. pombe)	RAD21	5885	NP_006256	C07000 IAIN	ŧ	200600 s at
ribosomal protein L37	RPL37	6167	NP 000988	NM 000997	ŧ	200607 s at
ribosomal protein S25	RPS25	6230	NP_001019	NM_001028	5 5	200091 S at
ribosomal protein S7	RPS7	6201	NP_001002	110100 MN	ŧ	200082 s at
IK cytokine, downregulator of HLA II	ķ	3550	NP_006074	NM_006083	40	200065 at
ribosomal protein L30	RPL30	6156	NP_000980	NM_000989	ŧ	200062 s at
ras homolog gene family, member A	RHOA	387	NP_001655	NM_001664	ŧ	200000 s at
nuclear DNAbinding protein	CID	10438	NP_006324	NM_006333	ŧ	2000050 s at
sperm associated antigen 7	SPAG7	9552	NP_004881	NM 004890	5 5	200002 at
zinc finger protein 146	ZNF146	7705	NP_009076	NM_007145	ŧ	200062 at
defender against cell death 1	DADI	1603	NP_001335	NM_001344	ŧ	200046 at
ATPbinding cassette, subfamily F (GCN20), member 1	ABCFI	23	NP_001081	NIM 001090	i é	m_c.ooo_
enhancer of rudimentary homolog (Drosophila)	ERH	2079	NP 004441	001000 MIN	6	200045 at
proteasome (prosome, macropain) subunit, beta type	PSMB2	0,000	14 _002/00	NIM 001450	à	200043 at
ribosomal protein S11	RPSII	0200	NF 001000	NM 00704	4	200039 s at
ribosomal protein L19	RPL19	6143	NP 000972	196000 MN	B E	200031 s at
ribosomal protein L34	RPL34	6164	NP_000986	MIN 000001	É	2000029 at
ribosomal protein L27	RPL27	6155	NP_0009/9	NIM 000005	A d	200026 at
ribosomal protein L18	RPL18	6141	NP_000970	NM 0009/9	ŧ	200022 at
ribosomal protein S13	RPS13	6207	NP 001008	/10100 IAN	ŧ	10 010002
ribosomal protein L24	RPL24	6152	NP 000977	NM_000986	ŧ	200013 at
ADPribosylation factor 3	ARF3	377	NP_001650	NM 001659	40	200011 s at
signal recognition particle 14kDa (homologous Alu	SRP14	6727	NP_003125	NN 003134	ŧ	200001_41
ribosomal protein L28	RPL28	8619	NP 000982	166000 Talvi	3 6	200007 of
ribosomal protein 1.35	RPL35	11224	NP 009140	NIM 000001	ð ð	200003 s at
hypothetical protein FLJ13197	FLJ13197	/966/	NP 0/8890	410470 Talvi	ài	200000 at
hypothetical protein FLJ31413	FLJ31413	100001	NID 079900	NM 02461	4D	1570414 x at
Ras homolog enriched in brain like 1	KHEBL1	121208	NID 660770	25521 WN	45	1570373 at
keratin 6L	KRT6L	338785	NP /8/028	NM 144502	5 6	1570253 a at
hypometical gene supported by BC039326				100201	3	1560000 01

	HEDAIA	2000	ND 005774	NM 005345	40	200/99 at
myeloid cell leukemia sequence (BCL2related)	MCLI	4170	NP 068779	NM 021900	ŧ	00700 A dt
myeloid cell leukemia sequence 1 (BCL2related)	MCLI	4170	NP 068//9	006170 MIN	3 8	200707 S at
myeloid cell leukemia sequence 1 (BCL2related)	MCLI	4170	NP 068//9	006170 IAIN	À d	200797 s at
phosphoprotein enriched in astrocytes 15	PEA 15	8682	NP 003/39	000 COO TAIN	40	200796 s at
ribosomal protein S15a	RPS15A	6210	010100 dN	MINI 001019	40	200788 s at
methionine adenosyltransferase II, alpha	MAT2A	4144	NP 005902	NW 001010	h 8	200781 s at
ribosomal protein, large, P1	RPLP1	6176	NP 000994	NM 001003	ŧ	200768 s at
calumenin	CALU	813	NP 001210	NM 001219	ŧ	200752 s at
ferritin, heavy polypeptide 1	FTHI	2495	NP_002023	NM 002032	ŧ	200746 S at
guanine nucleotide binding protein (G protein) be	GNBI	2782	NP_002065	NM 002074	ŧ	200744 s at
ribosomal protein S27 (metallopanstimulin 1)	RPS27	6232	NP_001021	NM_001030	ŧ	200741 3 41
nascentpolypeptideassociated complex alpha polypep	NACA	4666	NP_005585	NM_005594	5 5	200733 x at
ADPribosylation factor 3	ARF3	377	NP_001650	NM 001659	ŧ	200735 at
protein tyrosine phosphatase type IVA, member 1	PTP4A1	7803	NP_003454	NM_003463	ŧ	200733 S at
ARP2 actinrelated protein 2 homolog (veast)	ACTR2	10097	NP_005713	NM 005722	ŧ	200720 at
ribosomal protein L10	RPL10	6134	NP_006004	NM_006013	ŧ	200728 X at
ribosomal protein L13a	RPL13A	23521	NP 036555	NM 012423	ŧ	200735 x at
glutamicoxaloacetic transaminase 2 mitochondrial	GOT2	2806	NP_002071	NM_002080	45	200716 at
dynein, cytoplasmic, light polypentide I	DNCLI	8655	NP_003737	NM_003746	£	200709 at
NiemannPick disease, type C2	NPC2	10577	NP_006423	NM 006432	1	200701 at
KDEL (LysAspGluLeu) endoplasmic reticulum protein	KDELR2	11014	NP_006845	NIVI 000834	i é	
gelsolin (amyloidosis, Finnish type)	GSN	2934	NP_000168	NM 0001//	ŧ	200070 s at
ribosomal protein L32	RPL32	6161	NP_000985	NM 000994	t	200604 S at
spectrin, beta, nonerythrocytic 1	SPTBN1	6711	NP_003119	NM 003128	ŧ	200027 X at
ubiquitinconjugating enzyme E2D 3 (UBC4/5 homolog,	UBE2D3	7323	NP_003331	NM_003340	ŧ	200007.3 at
translocase of outer mitochondrial membrane 20 hom	TOMM20	9804	NP_05580	NMI_014/65	ŧ	200660 5 34
signal sequence receptor, beta (transloconassociat	SSR2	6746	NP 003136	NM 01476	à	200662 5 31
glutamateammonia ligase (glutamine synthase)	GLUL	2752	NP 002056	C90700 IAIN	3 8	200657 at
eukaryotic translation initiation factor 3. subuni	EIF3S8	8663	NP 003743	NM 003752	ŧ	200647 x at
superoxide dismutase I, soluble (amvotrophic later	1000				,	200617

actin binding LIM protein 1	ABLIMI	3983	NP_006711	NM_006720	4D	200965_s_at
ribosomal protein L31	RPL31	6160	NP 000984	NM 000993	40	200905 X at
clathrin, light polypeptide (Lca)	CLTA	1211	NP_009027	NM 007096		200960 x at
fusion (involved in t(12;16) in malignant liposarc	FUS	2521	NP 004951	NM 004960		18 6C6007
syndecan binding protein (syntenin)	SDCBP		NP 003616	C70C00 IAIN	5 5	200050 5 61
ribosomal protein S20	RPS20		NP_001014	520100 IAIN	1 1 5	200000 s at
arginineglutamic acid dipeptide (RE) repeats	RERE		NP 036234	NM 001000		200939 S at
ribosomal protein S23	RPS23		NP 001016	CZ0100 IMN		200000 a a
cytochrome c oxidase subunit VIa polypeptide I	COX6A1	1337	NP 004364	NM 004373	ŧ	200926 at
solute carrier family 3 (activators of dibasic and	SLC3A2		NP 002385	NM 002394	ŧ	200924 S at
polyhomeoticlike 2 (Drosophila)	PHC2	1912	NP 004418	NM_004427	ŧ	200021 11 616007
transgelin 2	TAGLN2	8407	NP 003555	NM 003564	45	200010 31
magnesiumdep					5	200016
protein phosphatase 1G (formerly 2C).	PPMIG	5496	NP_002698	NM_002707	4D	200913_at
ribosomal protein, large P2	RPLP2	1819	NP_000995	NM_001004	4D	200909 s at
major histocompatibility complex, class 1. E	HLAE	3133	NP_005507	NM_005516	40	200905 x at
splicing factor, arginine/serinerich 10 (transform	SFRS10	6434	NP_004584	NM_004593	40	200895 at
ribosomal protein L23	RPL23	9349	NP_000969	NM_000978	ŧ	200000 S at
creatine kinase, brain	CKB	1152	NP_001814	NM 001823	ŧ	200869 at
nucleolar protein 5A (56kDa with KKE/D repeat)	NOLSA	10528	NP_006383	NM_006392	6	2008/4 s at
RABIIA, member RAS oncogene family	RABIIA	8766	NP 004654	NM_004663	£	200865 S at
nuclear receptor corepressor 1	NCORI	9611	NP_006302	NM_006311	ŧ	200657 S at
guanine nucleotide binding protein (G protein), be	GNB2	2783	NP_005264	NM_005273	40	200857 x at
glutamylprolyltRNA synthetase	EPRS	2058	NP_004437	NM_004446	ŧ	200843 s at
ribosomal protein S21	RPS21	6227	NP_001015	NM_001024	40	200834 s at
procollagenlysine 1, 20xoglutarate 5dioxygenase 1	PLOD1	5351	NP 000293	NM_000302	ŧ	200827 at
small nuclear ribonucleoprotein D2 polypeptide 16.	SNRPD2	6633	NP_004588	NM_004597	ŧ	
ribosomal protein L29	RPL29	6159	NP 000983	NM 000992	ŧ	3C26007
ribosomal protein S15	RPS15	6209	NP 001009	810100 WN	ŧ	200822 s at
ATP synthase, H+ transporting, mitochondrial F1	ATP5O	539	NP_001688	NM_00109/	ŧ	200010 8
ribosomal protein S10	RPS10	6204	NP 001005	NM 001014	ŧ	200819 X at
heat shock 70kDa protein 1A	HSPAIA	3303	NP_005336	NM_005345	4D	200800 s at
					-	

	NEEDI O	4780	NP 006155	NM 006164	£	201146 at
HSI binding protein	HAXI	10456	NP_006109	NM 006118	ŧ	201145 at
major histocompatibility complex, class II, DP bet	HLADPB1	3115	NP 002112	NM_002121	ŧ	201137 S at
cytochrome c oxidase subunit VIIc	COX7C	1350	NP 001858	NM 001867	ŧ	201127 x at
heterogeneous nuclear ribonucleoprotein H2 (H")	HNRPH2	3188	NP_062543	NM_019597	ŧ	201137 St
splicing factor, arginine/serinerich 7, 35kDa	SFRS7	6432	NP_006267	NM 006276	ŧ	201122 at
cytochrome c oxidase subunit 8A (ubiquitous)	COX8A	1351	NP 004065	NM 004074	ŧ	201130 - at
carboxypeptidase E	CPE	1363	NP 001864	NM 001873	ŧ	201110 S at
CSE1 chromosome segregation [like (veast)	CSEIL	1434	NP_803185	NM_177436	ŧ	201117 at
thrombospondin I	THBSI	7057	NP 003237	NM_003246	ŧ	201111 S at
ribosomal protein \$29	RPS29	6235	NP_001023	NM_001032	\$	201094 at
dynactin I (n150 alived homolog Decombine)	DCTNI	1639	NP_004073	NM_004082	4D	201082 s at
general transcription factor II :	GTF2I	2969	NP_001509	NM_001518	4D	201065 s at
poly(A) binding protein cytoplasmic 4 (ind. ind	PABPC4	8761	NP_003810	NM_003819	4D	201064 s at
heterogeneous nuclear ribounclearests An	HNRPA0	10949	NP_006796	NM_006805	4D	201054 at
dual specificity phosphatase 1	DUSPI	1843	NP_004408	NM_004417	4D	201044 x at
guanine nucleotide hinding protein (C protein) of	GNAI2	2771	NP_002061	NM_002070	40	201040 at
L3hydroxyacylCoenzyme A dehydrogenase, short chain	HADHSC	3033	NF_005518	NIM DOOD Z	d	
eukaryotic translation initiation factor 5B	EIF5B	9669	NP 056988	MN 005327	3 8	201036 s at
destrin (actin depolymerizing factor)	DSTN	11034	NP 006861	0/8000 MM	ð	201024 v at
thioredoxin interacting protein	TXNIP	10628	NP_006463	NM 006472	à é	201022 s at
thioredoxin interacting protein	TXNIP	10628	NP 006463	NM 006472	ŧ	201010 s at
thioredoxin interacting protein	TXNIP	10628	NP_006463	NM 006472	ŧ	201000 S at
signal sequence receptor, delta (transloconassocia	SSR4	6748	NP_006271	NM 006280	ŧ	201008
alanyltRNA synthetase	AARS	16	NP_001596	C00100 INN	ŧ	201000 at
cytoskeletonassociated protein 4	CKAP4	10970	NP 006816	C78900 ININ	ŧ	201000 at
cytoskeletonassociated protein 4	CKAP4	10970	NP_006816	C78900 IMM	ŧ	2000000 5 41
hypoxiainducible factor 1, alpha subunit (basic he	HIFIA	3091	NP_001521	NM 001550	ð	200000 at
CD59 antigen p1820 (antigen identified by monoclon	CD39	700	141 000002			200000
pyruvate dehydrogenase (lipoamide) alpha 1	PUHAI	06100	C/7000 dN	NM 000611	40	200985 s at
transmembrane 4 superfamily member 8	LIVI4SES	5160	ND 00075	NM 000284	4D	200980 s at
uansinemorane 4 superfamily member 8	THATTO	10000	ND 005715	NM 005724	4D	200973 s at
1	- MAY	100	7.Centebra 1111			

coatomer protein complex, subunit beta	COLD	1010	111 001000			
THE CONTRACT (S. COLEVISIDE)	COBB		NP 057535	NM 016451	4D	201358 s at
YMEIlike I (S. cerevisiae)	YMEILI	10730	NP_055078	NM_014263	45	201552 at
glutathione peroxidase 3 (plasma)	GPX3	2878	NP_002075	NM 002084	ŧ	201353 at
adiponectin receptor 2	ADIPOR2	79602	NP 0/8827	1CC#20 IMM	t	201240 4
homolog,	086202	1022		NIA OOASSI	Ď	201346 at
stilati fluctear ribonucleoprotein polypeptide C	Diago.	7277	NP 003330	NM 003339	4D	201343 at
email nuclear sike and contex (white D I blue domain)	SNRPC		NP 003084	NM 003093	4D	201342 at
ectodermalneural cortex (with DTD112 4	ENCI	8507	NP_003624	NM_003633	4D	201341_at
vets erythroblastosis virus E26 oncogene homolog	ETS2	2114	NP_005230	NM_002239	ŧ	201020 gt
EBNA1 binding protein 2	EBNA1BP2	10969	NP 006815	NM 00522	ð	201328 at
serine/threonine kinase 25 (STE20 homolog, yeast)	S1 K25	10494	WF_000505	1111 0000774		20122
chromosome 5 open reading frame 13	C5orf13	9315	NP 004763	NW 00274	\$ 8	201314 at
DEAD (AspGluAlaAsp) box polypeptide 48	DDX48	9775	NP_05555	NM 014/40	ð	201310 s at
peptidylprolyl isomerase A (cyclophilin A)	PPIA	5478	NP 066953	001120 MM	àŧ	201303 24
proteasome (prosome, macropain) subunit, alpha typ	PSMAS	5686	NP_002/81	NIM_002/90	1	201203
red				00000	3	201274 at
aldoketo reductase family months B1 (11)	AKRIBI	231	NP 001619	NM_001628	4D	201272_at
ATPase Na+/K+ transporting bets 1 =-1	ATPIBI	481	NP_001668	NM_001677	4D	201242 s_at
capping protein (actin filament) muscle Zine, ap	CAPZA2	830	NP_006127	NM_006136	4D	201238 s at
capping protein (actin filament) muscle Zline ala	CAPZA2	830	NP_006127	NM_006136	4D	201237_at
BTG family, member 2	BTG2	7832	NP_006754	NM_006763	#	201232 S at
NADH dehydrogenasc (ubiquinone) 1 beta subcomplex,	NDUFB8	4714	NP_004995	NM_003004	d	201225
subcomplex,				2000	È	201227 8 9+
C)samin D (Stellin D)	NDIFER	4714	NP 004995	NM 005004	4D	201226_at
everatin B (classe B)	CSTB	1476	NP 000091	001000 MN	4D	201201_at
nhosnhatidylinosital transfer mustain -1-1	PITPNA	5306	NP_006215	NM 006224	4D	201192 s at
protease, serine 11 (IGE hinding)	PRSS11	5654	NP_002766	NM_002775	4D	201185 at
guanine nucleotide binding protein (G protein) al	GNA13	2773	NP_006487	NM_006496	4	201181 at
pumilio homolog 1 (Drosophila)	PUM1	9698	NP 055491	NM_014676	4D	201164 s at
NmvristovItransferase	ITMN	4836	NP_066565	NM_021079	45	201158 at
RAB5C, member RAS oncoone family	RAB5C	5878	NP_004574	NM 004583	46	201156 s at

proteasome (prosome macropain) subunit alaba	PSMA3	5684	NP_002779	NM_002788	4D	201532 at
ATPase, H+ transporting, lysosomal 14kDa, V1 subun	ATP6V1F	9296	NP_004222	NM_004231	ŧ	TD / 20102
ubiquitinconjugating enzyme E2N (UBC13 homolog, ye	UBE2N	7334	NP_003339	NM_003348	ŧ	201527 A at
small nuclear ribonucleoprotein polypeptide N	SNRPN	6638	NP_003088	NM 003097	ŧ	201524 × 21
translocase of outer mitochondrial membrane 70 hom	TOMM70A	9868	NP_055635	NM_014820	ŧ	10 C10102
nuclear cap binding protein subunit 2, 20kDa	NCBP2	22916	NP 031388	NM 007362	ŧ	201517_dt
protein phosphatase 1, regulatory (inhibitor) subu	PPPIRII	6992	NP 068778	NM 021959	ŧ	201517 S at
ribosomal protein L41	RPL41	6171	NP_066927	NM_021104	ŧ	201500 - at
AHAI, activator of heat shock 90kDa protein ATPase	AHSAI	10598	NP_036243	NWI OLZIII	đ	201402
peptidylprolyl isomerase F (cyclophilin F)	PPIF	10105	NP 005720	NM 005729	ŧ	201490 S at
quiescin Q6	QSCN6	5768	NP_002817	NM_002826	ŧ	201400 at
sequestosome	SQSTMI	8878	NP_003891	NM_003900	8	2014/1_s_at
NAD(P)H dehydrogenase, quinone I	NQ01	1728	NP_000894	NM_000903	45	201400 S at
Ras homolog enriched in brain	RHEB	6009	NP_005605	NM 005614	ŧ	201469 x at
dihydropyrimidinaselike 3	DPYSL3	1809	NP_001378	NM_001387	ŧ	201453 S at
ribosomal protein L37a	RPL37A	6168	NP_000989	NM_000998	ŧ	201429 S at
methylosome protein 50	MEP50	79084	NP_077007	NM 024102	ŧ	201420 at
SRY (sex determining region Y)box 4	SOX4	6659	NP_003098	NM_003107	ŧ	201421 at
SRY (sex determining region Y)box 4	SOX4	6659	NP_003098	NW_003107	ŧ	201417 at
hydroxysteroid (17beta) dehydrogenase 4	HSD17B4	3295	NP_000405	NM_000414	ŧ	301416 ot
low density lipoprotein receptorrelated protein 10	LRP10	26020	NP 054764	NM_014045	ŧ	201412 at
ribosomal protein L36a	RPL36A	6173	NP_066357	NM 021029	ŧ	201406 at
adrenergic, beta, receptor kinase 1	ADRBK1	156	NP_001610	NM_001619	8	201401 s at
proteasome (prosome, macropain) subunit, beta	PSMB3	5691	NP_002786	NM_002/95	ć	-00-100_at
insulinlike growth factor 2 receptor	IGF2R	3482	NP 000867	9/8000 MIN	ð	201372 3 at
neighbor of BRCA1 gene 1	NBRI	4077	NP 005890	NM 005899	ŧ	201204 S at
ubiquitin associated protein 2like	UBAP2L	9898	NP_055662	NM 014847	É	201281 at
ornithine decarboxylase antizyme 2	OAZ2	4947	NP 002528	/ CC200 IMN	ŧ	201377 of
influenza virus NSIA binding protein	IVNSLABP	10625	NP_006460	NM_006469	÷	
coatomer protein complex subunit beta	COPB	1313	141 00000			200

nucleoside phosphorylase	NP	4860	NP_000261	NM 000270	t	201092 S at
apoptosis inhibitor 5	API5	8539	NP 006586			
chromosome 14 open reading frame 92	C14orf92		NP 055645			201687 s at
peptidase (mitochondrial processing) beta	PMPCB	9512	NP_004270			
arsenate resistance protein ARS2	ARS2	51593	NP 056992			a
DC12 protein	DC12		NP_064572			
ribosomal protein S17	RPS17	6218	NP 001012			i ii
ADPribosylation factorlike 1	ARLI	400	NP 001168			
interferon gamma receptor 2 (interferon gamma tran	IFNGR2	3460	NP_005525			9
bone marrow stromal cell antigen 2	BST2	684	NP_004326			
acid phosphatase 1, soluble	ACPI	52	NP_004291			12
aspartyltRNA synthetase	DARS	1615	NP_001340			
staphylococcal nuclease domain containing 1	SNDI	27044	NP_055205			
aldehyde dehydrogenase 9 family, member A1	ALDH9A1	223	NP_000687			
nuclear phosphoprotein similar to S. cerevisiae PW	PWPI	11137	NP_008993			9
repressor of estrogen receptor activity	REA	11331	NP 009204			
ornithine aminotransferase (gyrate atrophy)	OAT	4942	NP 000265			
keratin 18	KRT18	3875	C17000 AN			4
protein phosphatase 4, regulatory subunit 1	PPP4R1	9989	NE 000120			
response,			ND 005125	NM 005124	4D	201594 s at
likely ortholog of mouse immediate early	LEREPO4	55854	NP_060941	NM_018471	ŧ	201070_S_dt
eukaryotic translation initiation factor 3, subuni	EIF3S3	8667	NP_003747	NM_003756		
splicing factor proline/glutamine rich (polypyrimi	SFPQ	6421	NP 005057	000000 MM		1
Sec23 homolog B (S. cerevisiae)	SEC23B	10483	NP_006354	NM 006565		
hypothetical protein DJ971N18.2	DJ971N18.2	56255	NP_066979	NM_021126		12
nonmetastatic cells 1, protein (NM23A) expressed	NME	4830	NP_000260	Min 000209		
dCMP deaminase	DCTD	1635	NP 001912	176100 MIN	3 6	201577 at
low molecular mass ubiquinonebinding protein (9.5k	QPC	27089	NF_05217	NM 00102		-
dual specificity phosphatase 3 (vaccinia virus pho	DUSP3	1845	NP 004081	NM 004090		201568 24
ubiquitinlike 3	UBL3	5412	NP 009037	NM 007106		201325 at
ţyp						303100

		200	NID 005700	NM 005808	4	201407 6 91
interleukin 13 receptor, alpha 1	IL13RA1 ii	3597	NP_001551	000 INN	ŧ	10
interleukin 13 receptor, alpha 1	IL13RA1 i	3597	NP_001551	NM 001560	ŧ	201067 at
paraoxonase 2	PON2	5445	NP_000296	NM 000303	ŧ	2010/0 81
myelin protein zerolike	MPZL1 r	9019	NP 003944	NM 003953	ŧ	2010/4 at
ORF	LOC51035	51035	NP 056937	NVI 015853	5	201071 3 dt
transducin (beta)like 1Xlinked	TBL1X t	6907	NP_005638	NM_005647	ŧ	201868 S at
leucine rich repeat (in FLII) interacting protein	LRRFIPI	9208	NP_004726	NM_004/35	ŧ	2010/0 S at
proteoglycan I, secretory granule	PRGI	5552	NP_002718	NM 002/2/	ŧ	201027 at
cell division cycle 25B	CDC25B	994	NP_068659	NM 021873	ŧ	201050 ct
neural precursor cell expressed, developmentally d		4738	NP_006147	NM_006156	ŧ	201840 at
tumorassociated calcium sional transducer	Ξ	4072	NP 002345	NM_002354	4D	201839 s at
protein kinase. AMPactivated beta I noncaralytic	ıBI	5564	NP_006244	NM_006253	4D	201835 s at
neuroepithelial cell transforming some		10276	NP_005854	NM_005863	4D	201830 s at
neuroenithelial cell transforming game l		10276	NP_005854	NM_005863	4D	201829_at
ring finger protein 14	4	9604	NP_004281	NM_004290	4D	201824_at
endoglin (OslerRend) Weber syndrome 1)		2022	NP_000109	NM_000118	4D	201809 s at
valvitRNA synthetase 2	S2	7407	NP_006286	NM_006295	4D	201797_s_at
lamin B recentor		3930	NP_002287	NM_002296	4D	201795_at
vrel reticuloendotheliosis viral oncorrene homological		5970	NP_068810	NM_021975	45	201783 s at
ring finger protein 13		11342	NP_009213	NM_007282	4D	201780 s at
KIAA0494 gene product	4	9813	NP_055589	NM_014774	4D	201778 s at
secretory carrier membrane protein 3	3	10067	NP_005689	NM_005698	4Đ	201771_at
small nuclear ribonic leoprotein polymentals A		6626	NP 004587	NM_004596	4D	201770_at
cytochrome c oxidase subunit VIc	SC	1345	NP 004365	NM_004374	40	201754 at
tumor protein p53 (LiFraumeni syndrome)	TP53	7157	NP_000537	NM_000546	4b	201746_at
NADH dehydrogenase (ubiquinone) FeS protein 3 30k	NDUFS3	4722	NP_004542	NWI_004551	ŧ	201770_at
membraneassociated ring finger (C3HC4) 6	MARCH6	10299	NP 005876	NM 005885	à ê	201737 S at
UDPNacetylalphaDgalactosamine:polypeptide Nacetylg	GALNTI	2589	NP_065207	1414 020474	i	
erythrocyte membrane protein band 4.11ike 2	EPB41L2	2037	NP 001422	NM 030434	å é	201773 s at
cyclin D3	CCND3	896	NP 001751	NM_001760	ŧ	201719 at
DNA (cytosine5)methyltransferase 1	LIMINIC	1,00				201200

peptidase D cytochronne c oxidase subunit VIIb	COX7B cy	1349	ND 000007		dD Ch	707 P
ptidase D						
0		2104				202110 at
bromodomain containing 4		1				202108 at
		22/76	NP 055114	NM 014299	4D	202102 s at
vral simian leukemia viral oncogene homolog B	RALB vral	5899	NP_002872	NM_002881	+5	202101 S at
vral simian leukemia viral oncogene homolog B	RALB vrai					202101 6 64
HMT1 hnRNP methyltransferaselike 1 (S. cerevisiae)	HRMT1L1 H	32/3	NP_001526			00100 0
OGT(OGleNAc transferase)interacting protein 106 KD		22906	NP_055780		5 6	202098 s at
NADH dehydrogenase (ubiquinone) 1, alpha/beta subc	NDUFABI N	4706	NP_004994			10 0202 at
low density lipoprotein receptor (familial hyperch	LDLR lo	3949	NP_000518			202006_S_at
sel1 suppressor of lin12like (C. elegans)	SELIL se	6400	NP_005056			202069 S at
sell suppressor of lin12like (C. elegans)	SEL1L se	6400	NP_005056			202061_s_at
TGFB1induced antiapoptotic factor 1	TIAF1 T	9220	NP 004731			702027 at
RB linducible coiledcoil 1	RB1CC1 R	9821	NP_055596			202034 X at
ribosomal protein L38	RPL38 ri	6169	NP_000990			202029 x at
putative translation initiation factor	SUI1 pi	10209	NP_005792			10 V 170707
LanC lantibiotic synthetase component Clike 1 (bac	LANCLI L	10314	NF_006046			202021
acetylCoenzyme A acyltransferase 2 (mitochondrial	ACAA2 au	10449	NF_006102			202020 s at
Ras suppressor protein 1	RSUI R	6251	NP_036557	NM 012425		202003 s at
				NM_021122		201963_at
decay accelerating factor for complement (CD55,	DAF d	1604	NP_000565	NM_0005/4		-01727 s_at
solute carrier family 20 (phosphate transporter),	SLC20A1 s	6574	NP 005406	NM 003413		201025 G at
hypothetical protein FLJ10618	FLJ10618 h	55186	NP 060625	0018105 MM_018105		10 C 116107
G1 to S phase transition 1	GSPT1 C	2935	NP 002085	NM 002094		201017 3 01
FERM, RhoGEF (ARHGEF) and pleckstrin domain protei	FARP1 F	10160	NP_005/5/	NM 00000		201012 - 24
II, po			ND OCCUPANT	225500 WIN	45	201910 at

monitorios)					-	
ubiquitinconjugating enzyme E2B (RAD6	UBE2B	7320	NP_003328	NM_003337	ŧ	202334_S_at
polycystic kidney disease 1 (autosomal dominant)	PKDI	5310	NP_000287	NM 000296	ŧ	202324 S at
HLAB associated transcript 8	BA18	ľ	NF 000/00	MIN 000707	à	10 3 8CECOC
geranylgeranyl diphosphate synthase I			NE_004020	NM 00700	4	202326 at
G (KNAN) II (DINA directed) polypeptide			ND OO 1000	NM 00/837	45	202322 s at
nolymense (DNA) II (DNA dimense)	POLROG	5436	NP 002687	NM_002696	4D	202306_at
NADH dehydrogenase (ubiquinone) I alpha subcomplex	NDUFA1	4694	NP_004532	NM_004341	ŧ	0.770 at
RER1 retention in endoplasmic reticulum 1 homolog		9	NP_008964	NM 004841	à é	202208 at
laminin, gamma 2	LAMC2	3918	NP 005555	70CC00 MINI	ð	202205 04
TRAF and TNF receptor associated protein	TTRAP	51567	NP_057698	NM 005572	ð	202200 at
translocase of outer mitochondrial membrane 40 hom	I OMM40	10407	000105			20200
transcription factorlike 1		6944	NP 005988	NM 006114	5 8	202264 s at
phosphonoformate immunoassociated protein 5	PFAAP5	10443	NP 055702	NM_014887	ŧ	202239 S at
cyclindependent kinase 4	CDK4	1019	NP_000066	NM_000075	ŧ	202246 S at
nicotinamide Nmethyltransferase	TMNN	4837	NP_006160	NM_006169	ŧ	20224/ at
stromal cell derived factor receptor 1	SDFRI	27020	NP_059429	NM_017455	40	202228 s at
KIAA0907 protein	KIAA0907	22889	NP_055764	NM_014949	ŧ	202220_at
LSM3 homolog, U6 small nuclear RNA associated (S.	LSM3	27258	NF_055/8	14400 1410 Talki	1	
autocrine motility factor receptor	AMFR	267	NP_001135	NM 01144	ð	202209 st
myotubularin related protein 3	MTMR3	8897	NP 066576	NM 021090	ŧ	202176 S at
L1M domain kinase 2	LIMK2	3985	NP_005560	NM_005569	ŧ	202100 at
protein phosphatase 2, regulatory subunit B (B56)	PPP2R5A	5525	NP_006234	NM_006243	40	202187 s at
TAF9 RNA polymerase II, TATA box binding protein (TAF9	6880	NP_003178	NM 003187	ŧ	200 at
tubulin, beta 3	TUBB3	10381	NP 006077	NM 006086	ŧ	202134 X at
neural precursor cell expressed, developmentally d	NEDD9	4739	NP 006394	NM_006403	40	202149 at
adenviosuccinate ivase	ADSL	158	NP_000017	NM_000026	4D	202144 s at
ger. MYND domain containing	ZMYNDII	10771	NP_006615	NM 006624	4D	202136_at
vabl Abelson murine leukemia viral oncogene homolo	ABLI	25	NP_005148	NM_005157	ŧ	202123_s_at

FXYD domain containing ion transport regulator 3	FXYD3	3349	706C00 JAI	1117 000 111		
and protein i		53.10	ND 005062	NM 005971	4D	202488 s at
RAN binding protein 1	RANBPI	5902	NP 002873	NM_002882	4D	202483_s_at
27.7				NM_006326	4D	202475_at
catenin (cadherinassociated protein) alphalika I	CTNNALI	8727	NP_003789	NM 003798	4D	202468 s at
6phosphofructo2kinase/fructose2,6biphosphatase	PFKFB3	5209	NP_004557	NM_004566	ŧ	202404 S at
lipin 2	LPIN2	9663	NP 055461	NM_014646	ŧ	202464 S at
chromosome 9 open reading frame 60	C9orf60	10444	NP 006327	NM_006336	ŧ	202450 s at
retinoid X receptor, alpha	RXRA	6256	NP_002948	VM 00295/	ŧ	20245/ 20245/ 20245/
SPFH domain family, member 1	SPFHI	10613	NP_006450	NM_006459	ŧ	202444 S at
adaptorrelated protein complex 3, sigma 1 subunit	AP3S1	1176	NP_001275	NM_001284	ŧ	202444 at
cytochrome P450, family 1, subfamily B, polypeptid	CYPIBI	1545	NP_000095	NW 000104	ŧ	202424 S at
vmyc myelocytomatosis viral oncogenc homolog (avia	MYC	4609	NP_002458	NM_00246/	ŧ	202434 s. at
diazepam binding inhibitor (GABA receptor modulato	DBI	1622	NP 065458	NM_020348	ě	202421
DKFZP564B167 protein	DKFZP564B167	25874	NP_056230	NM 030518	ŧ	202427 S at
retinoid X receptor, alpha	RXRA	6256	NP_002948	NM_002957	ŧ	202427 S at
acylCoA synthetase longchain family member 4	ACSL4	2182	NP_075266	NM_022977	#	202426 at
Yip1 interacting factor homolog (S. cerevisiae)	YIFI	10897	NP_065203	NM 020470	6	202420 at
PRP31 premRNA processing factor 31 homolog (yeast)	PKPF31	20121	144 144 144	.an_015025	i	200
cysteinyltRNA synthetase	CARS	833	NP 001/42	107100 IAIN	4) 6	202408 s at
adaptorrelated protein complex 3, sigma 2 subunit	AP3S2	10239	NP 005820	NM 001751	4 5	202377 S at
transcription elongation regulator 1	TCERGI	10915	NP_006697	NM_006/06	ŧ	202390 at
ATPbinding cassette, subfamily F (GCN20), member 3	ABCF3	55324	NP_060828	MM 00 630 9	ě	20200 at
phosphatidylserine decarboxylase	PISD	23761	NP 055153	NM 014338	3 8	202394 s at
leptin receptor overlapping transcript	LEPROT	54741	NP 059996	NM 01/526	ŧ	202377 dt
transcription elongation factor A (SII)like 4	TCEAL4	79921	NP_079139	NM_024863	ŧ	2023/1 at
torsin family I, member A (torsin A)	TORIA	1861	NP 000104	NM 000113	ŧ	202371 at
fatty acid binding protein 5 (psoriasisassociated)	FABP5	2171	NP 001435	NM_001444	ŧ	202345 s at
ubiquitinconjugating enzyme E2B (RAD6 homolog)	UBE2B	7320	NP_003328	NM_003337	ŧ	202303 S AL
				1	5	202226

membraneassociated ring finger (C2HCA) 7	MARCH7	64844	NP_073737	NM_022826	40	202034 X at
lysophosphatidylglycerol acyltransferase 1	LPGATI	9926	NP_055688	NM_014873	ŧ	202651 at
ribosomal protein S19	RPS19	6223	NP_001013	NM 001022	ŧ	202651 at 020202
tumor necrosis factor, alphainduced protein 3	TNFAIP3	7128	NP_006281	NIN 000290	ŧ	202047 3 41
tumor necrosis factor, alphainduced protein 3		7128	NP 006281	NM 006290	1 5	202042 s at
ADPribosylation factorlike 3	ARL3		NP_004302	NM 002200	ð	202041 81
polymerase (RNA) II (DNA directed) polypeptide K,	POLR2K	0440	NF_003023	NA 001311	9 6	202641 at
amyloid beta precursor protein (cytoplasmic tail)		10013	NID 005035	NM 005034	4	202635 s at
chromosome 14 open reading frame 11		55837	NP 000923	NW 008280	4D	202630 at
procollagenlysine, 20x0glutarate 5dioxygenase 2	PLOD2	5352	NP_000926	NM 018453	4) 5	202020 s at
methyl CpG binding protein 2 (Rett syndrome)	MECP2	4204	NP 004983	NM 004992	ŧ	18 S 910707
cofactor required for Sp1 transcriptional activati	CRSP2	9282	NP_004220	NM 004229	ŧ	18 S 110707
nuclear receptor interacting protein 1	NRIPI	8204	NP_003480	NM_003489	ŧ	202600 s at
membrane interacting protein of RGS16	MIR16	51573	NP_057725	NM_016641	ŧ	202293 S at
singlestranded DNA binding protein 1	SSBPI	6742	NP_003134	NM 003143	ŧ	202503 s at
heat shock 70kDa protein 1B	HSPAIB	3304	NP_005337	NM 005346	ŧ	202501 at
DEAD (AspGluAlaAs) box polypeptide 19like	DDX19L	55308	NP_060802	NM_018332	ŧ	202501 - 10202
small nuclear ribonucleoprotein D3 polypeptide 18k	SNRPD3	0634	NP_004166	NW_004173	i d	200257
supervillin	SVIL	6840	NP 003165	NM 003175	ð	202567 at
DKFZP547E1010 protein	DKFZP547E101	26097	NP_056422	NW_01000/	ŧ	202200 a m
GCIPinteracting protein p29	P29	25949	NP_056299	NM 015484	ŧ	202505 s at
3hydroxy3methylglutarylCoenzyme A reductase	HMGCR	3156	NP_000850	NM_000859	ŧ	202540 S at
DKFZP564O123 protein	DKFZP5640123	25978	NP 054762	NM_014043	ŧ	202536 at
interferon regulatory factor 1	IRFI	3659	NP_002189	NM_002198	40	202531_at
CCCTCbinding factor (zinc finger protein)	CTCF	10664	NP_006556	NM_006565	40	202521_at
mutL homolog 1, colon cancer, nonpolyposis type	MLHI	4292	NP_000240	NM_000249	ŧ	202020 S BI
protein phosphatase 2, regulatory subunit B (B56)	PPP2R5D	5528	NP_006236	NM_006245	40	202515 s at
tumor necrosis factor, alphainduced protein ?	TNFAIP2	7127	NP 006282	NM_006291	4D	202509 s at
acylCoenzyme A dehydrogenase, C4 to C12 straight c	ACADM	34	NP_000007	NM_000016	8	202502_at
solute carrier family 2 (facilitated of neoce trans	SLC2A3	6515	NP_008862	INM 000951	t	200000000000000000000000000000000000000

	DRGI	4733	NP 004138	NM 004147	40	707010 at
drebrin I	DBNI	1627	NP 004386	NM 004395	ŧ	019C00
solute carrier family 1 (glial high affinity gluta	53		NP 004163	7/11400 TAIN	3 8	202000 at
phospholipase C, gamma 1			NP 8//963	110701 IAINI	ð	202800 at
mitogenactivated protein kinaseactivated protein k	MAPKAPK3	/86/	NF 004020	THOCOL MAN	4) 6	
NADH dehydrogenase (ubiquinone) I alpha subcomplex		4701	NP_004992	NW 003635	45 8	202787 s at
skeletal muscle and kidney enriched inositol phosp		51763	NP_057616	NM 005001	ð	202785 at
soc2 suppressor of clear homolog (C. elegans)	SHOC2	8036	NP_031399	NM 01/5/3	3 6	202777 at
splicing factor, arginine/serinerich 8 (suppressor	SFRS8	6433	NP_004583	NM 004592	à É	2027/3 s at
cyclin G2	12	901	NP 004345	NM_004354	40	2027/0 s at
acid phosphatase 2. vsosoma	ACP2	53	NP_001601	NM_001610	40	202/6/_at
fibrillin I (Marfan syndrome)		2200	NP_000129	NM_000138	4D	202766 s at
PALM2AKAP2 protein	KAP2	445815	NP_009134	NM_007203	4D	202759_s_at
regulatory factor Xassociated ankyrincontaining	RFXANK	8625	NP_003712	NM_003721	£	202738 s at
R3H domain (binds singlestranded nucleic acids) co	R3HDM	23518	NP_056176	NWI_015361	ŧ	202757_86
tryptophan rich basic protein	WRB	7485	NP_004618	NW 00402/	ŧ	202754 ot
guanylate binding protein 2, interferoninducible	GBP2	2634	NP 004111	021400 TAINI	à	202749 at
integral membrane protein 2A	ITM2A	9452	NP 004858	100+00 MINI	3 8	202749 24
latent transforming growth factor beta binding pro	LTBPI	4052	NP 000618	720000 MIN	4 6	202747 s at
ligase I, DNA, ATPdependent		3978	NP 000225	PCZ000 IAIN	4 6	202728 s at
BET1 homolog (S. cerevisiae)	BETI	10282	NP 003839	00000 MIN	È	202726 at
cytochrome c oxidase subunit IV isoform 1	=	1327	NP 001852	NM 005060	4 6	202710 at
serine/threonine kinase 17a (apoptosisinducing)	STK17A	9263	NP 004/51	NIV 001871	100	202608 v at
upstream binding transcription factor, RNA polymer	OBIT	/343	055040	200720	È	16 ° 509C0C
small nuclear ribonucleoprotein D1 polypeptide 16k	SNRPDI	6632	NF 000009	NIM DIADES	45 i	202692 s at
general transcription factor IIA, 2, 12kDa	G1F2A2	2958	ND 000000	NM 00402	4D	202690 s at
WiskottAldrich syndrome protein interacting protei		7456	NP_003378	NIM 00/402	4 6	202678 at
WiskottAldrich syndrome protein interacting protei	WASPIP	7456	NP_0033/8	NIM DOSSES	3 6	202665 s at
peroxisomal biogenesis factor 11B	LEVILD		17	200000	3	202663 at

vyes1 Yamaguchi sarcoma viral oncogene homolog 1			TD OOOCC	C9C000 MN	+	202943 s at
	YESI	7525	NP_005424	NI 005+55	10	
neuroblastomaamplified protein	NAG	51594	NP 056993	NM 005433	3 8	202933 s at
glutamatecysteine ligase, catalytic subunit	GCLC	2729	NP 001489	NM 001498	ŧ	202020 ot
ankyrin 2, neuronal	ANK2	287	NP_001139	NM 001148) t	202921 S at
CD97 antigen	CD97	976	NP_001775	NM 001784	ŧ	202921 at
Nijmegen breakage syndrome 1 (nibrin)	NBSI	4683	NP 002476	NM 002485	ŧ	202010 s at
Nijmegen breakage syndrome (nibrin)	NBSI	4683	NP 002476	NM_002485	3 5	000007 % at
Nijmegen breakage syndrome 1 (nibrin)	NBSI	4683	NP_002476	NM 002485	100	202902 x at
cathepsin S	CTSS	1520	NP_004070	NM_004079	ŧ	202905 s at
Freehommer, monteceptor type 800				NM_004648	40	202896_s_at
protein tyrosine phosphatase population (con arrange)	PTPNSI	140885	NP_542970	NM_080792	4D	202895 s at
CDC33 (cell division such 32 sect 1	CDC23	8697	NP_004652	NM_004661	4D	202892 at
microtubuleassociated protein 7	MAP7	9053	NP_003971	086500 WN	ŧ	702002 A 41
alanyl (membrane) aminopeptidase (aminopeptidase N	ANPEP	290	NP_001141	NM 600000	3 6	202860
DNAdamageinducible transcript 4	DDIT4	54541	NP 061931	80610 IAIN	45 8	202888 s at
nucleolar protein 7, 27kDa	NOL7	51406	NP_057251	NM_01616/	ŧ	20200Z A at
DnaJ (Hsp40) homolog, subfamily B, member 12	DNAJB12	54788	NP_001002762	NM 001002/62	ŧ	202820 at
transmembrane protein 4	TMEM4	10330	NP 055070	CC7110 IMM	è	20262 at
solute carrier family 16 (monocarboxylic acid tran	SLC16A3	9123	NP_004198	NW 004207	è	202020 o m
solute carrier family 16 (monocarboxylic acid tran	SLC16A3	9123	NP 004198	NM 004207	ÈÉ	202856 s at
phosphatidylinositol glycan, class C	PIGC	5279	NP 002633	NM_002642	ŧ	358COC
NADH dehydrogenase (ubiquinone) I beta subcomplex,	NDUFB7	4713	NP_004137	140 TAN TAN	ð	202016
fucosidase, alphaL 1, tissue	FUCAI	2517	NP_000138	NM 000147	ð 8	202839 s at
serine (or cysteine) proteinase inhibitor, clade A	SERPINAI	5265	NP_000286	C67000 IAIN	à	202033 of
GRIP and coiledcoil domain containing 2	GCC2	9648	NP 055450	NM 014635	3 8	202833 8 94
LIM domain containing preferred translocation part	LPP	4026	MP_000099	NM_003576	3 6	707837 at
transcription elongation factor B (SIII), polypept	TCEB3	6924	NP 00550	NIM 005579	45 6	202822 at
synovial sarcoma translocation, chromosome 18	SS18	0/60	NP 003160	NM 003108	40	202818 s at
TAR (HIV) RNA binding protein 1	TARBPI	6894	NP 005636	NM 005637	45	202817 s at
STAM binding protein	STAMBP	1001/	10 000 CA	NIM 005646		202813 at

SET domain, bifurcated 1	SETDB1	9869	NP 056564	204210 10432		
KIAA0040 gene product	KIAA0040	90/4	ND 035551	01000 MIN		
TATA box binding protein	IBP	0000	ND 055471	NM 014656		at
Secol Dela Subunit	TDD	8009	NP 003185	NM 003194	4D	203135_at
Social but the distribution of the second but the s	SECKIB	10952	NP 006799	NM_006808	4D	203133_at
inosito (cara) (cara)	IMPA2	3613	NP 055029	NM_014214	4D	203126_at
ubiquitin enerific protects 50	USP52	9924	NP_055686	NM_014871		at
2 John Physical Highlandin Pacifol 7				NM_014247	40	20309/ S at
mitochondrial translational initiation forces	MT1F2	4528	NP 002444	NM_002453		203095 at
protease serine 25	PRSS25	27429	NP_037379	NM_013247		202002 at
component of oligomeric goldi complex 2	COG2	22796	NP_031383	NM_007357		203073 at
kelchlike 21 (Drosombile)	KLHL21	9903	NP_055666	NM_014851		203058 at
				NM_014863		203000 at
forkhead box K2	FOXK2	3607	NP_004505	NM_004514		203064 s at
3'phosphoadenosine s'phosphoaniae synnase 2	PAPSS2	9060	NP_004661	NM_004670	40	203060 s at
3'phosphoadenosine 5'phosphoadence 2	PAPSS2	9060	NP_004661	NM_004670		203056 S at
breast carcinoma amplified (concess)	BCAS2	10286	NP_005863	NM_005872	40	203053 at
timeless homolog (Drosophila)	TIMELESS	8914	NP 003911	NM_003920	ŧ	202052 30 S at
ninjurin l	ILNIN	4814	NP_004139	NM 004148	5 6	203045 at
lysosomalassociated membrane protein 2	LAMP2	3920	NP_002285	NM 002294	ŧ	202045 at
erythrop						2020/2
uroporphyrinogen III synthase (corrected	UROS	7390	NP_000366	NM_000375	4D	203031_s_at
profein typesine phoenhates, receptor type, N pol	PTPRN2	5799	NP_002838	NM_002847	4D	203030 s at
protein tyrosine shooshotoo	PTPRN2	5799	NP_002838	NM_002847	4D	203029 s at
Tine finger and BTB description	ZBTBS	9925	NP_055687	NM_014872	4D	203026_at
hynothetical protein Henciting brote	HSPCIII	51491	NP_057475	NM_016391	4D	203023_at
Synovial sarcoma V baseline 2	SSX2IP	117178	NP_054740	NM_014021	4D	203019 x at
ribosoma protein 122	RPL23A	6147	NP_000975	NM_000984	4D	203012_x_at
membrane protein policitoral and regulat	MPPI	4354	NP_002427	NM_002436	4D	2029/4 at
dualspecificity tyrosing(x)phospholylation regular	DYRK2	8445	NP_006473	NM_006482	40	2029/1 s at
dualspecificity tyrosing(V) who exhaust as	DYRK2	8445	NP 006473	NM_006482	45	202958 s at
calnain 6	CAPN6	827	NP_055104	NM_014289	40	202966 at
regulatory factor Y \$ (influence Till A	RFX5	5993	NP_000440	NM 000449	4D	202963 at
ATP synthase, H+ transporting, mitochondrial F0	ATP5J2	9551	NP_004880	NM_004889	£	-02701_S_dt
						20200

LIDBNG-1-1-1-1	GAI NT3	1050	NP 004473	NM 004482	40	20009/ S at
kinesin family member 3C	KIF3C	3797	NP 002245	INIM 002234	ŧ	202207 5 41
kinesin family member 3C	KIF3C	3797	NP_002245	12C00 14M	3 6	13300 6 94
diacylglycerol kinase, alpha 80kDa	DGKA	1000	INF_001330	CECTOO TAIN	45	203389 at
cell division cycle 40 homolog (yeast)	CDC40	21362	NE 030973	NIN 01304	4D	203385 at
tripeptidyl peptidase II	TPP2	/1/4	707500 IN	NW 015801	4D	203376 at
tripeptidyl peptidase II	TPP2	2174	10 002002	190500 MN	4D	203375 s at
citiye binding protein	TOPO	7174	NP 003282	NM 003291	4D	203374 s at
emidicer of zeste nomolog 2 (Drosophila)	MVCDD	26292	NP 036465	NM 012333	4D	203361_s_at
suounit	EZHO	2146	NP 004447	NM 004456	4D	203358_s_at
adaptorrelated protein complex I, gamma I	APIGI	164	NP_001119	NM_001128	ŧ	202020_at
caveolin 2	CAV2	858	NP_001224	NM_001233	ŧ	202250 A
zinc finger protein 148 (pHZ52)	ZNF148	7707	NP_068799	NM 021964	ŧ	203319 s at
small nuclear ribonucleoprotein polynestide E	SNRPE	6635	NP_003085	NM_003094	40	203316 s at
GTP binding protein 6 (putative)	GTPBP6	8225	NP_036359	NM_012227	ŧ	203314 81
syntaxin binding protein 3	STXBP3	6814	NP_009200	NM_007269	1	203314 at
Jumonji, AT rich interactive domain 2	JARID2	3720	NP_004964	NM_004973	ŧ	202210 S at
vacuolar protein sorting 11 (yeast)	VPSII	55823	NP 068575	MM 021729	3 8	02200 0 00
DNA fragmentation factor, 45kDa, alpha polypeptide	DFFA	10/0	NE_004392	NIM 001700	3	203292 s at
chromosome 6 open reading frame 74	Cbort/4	07010	141/CO 1M	NM 004401	4D	203277 at
Fc fragment of IgG binding protein	FCOBP	0037	ND 057147	NM 016063	45	203259 s at
sarcoma amplified sequence	SAS	0302	NP 003772	NM 003890	\$	203240 at
adenine phosphoribosyltransferase	21 81	2000	ND OOSO73	NM 005981	4D	203227 s at
HIVI rev binding protein 2	ADDT	353	NP 000476	NM 000485	4D	203219_s_at
menymansier	HRR7	11103	NP 008974	NM 007043	4D	203202_at
5methyltetrahydrofolatehomocysteine	MTRR	4552	NP_002445	NM_002454	4D	203199_s_at
				NM_003138	40	203182 s at
glycine amidinotransferase (Larginine:glycine amid	GATM	2628	NP_001473	NM_001482	ŧ	2001/0-00
esophageal cancer associated protein	MGC16824	57020	NP 064710	NM 020314	ŧ	203178 at
KIAA0409 protein	KIAA0409	23378	NP_056139	NM_015324	ŧ	2021/1 S at
cAMP responsive element binding proteinlike 1	CREBLI	1388	NP 004372	NM 004381	ŧ	203100 at
tissue inhibitor of metalloproteinase 2	TIMP2	7077	NP_003246	NM_003255	40	20316/ at
A Milase (PKNA) anchor profess						

zinc fingers and homeoboxes 2	ZHX2	22882	NP_055758	NM_014943	10	-00000 at
CD4 antigen (p55)	CD4	920				4
calcium modulating ligand	CAMLG					
LSMI homolog, U6 small nuclear RNA associated (S.	LIMCT	,				
adenomatosis polyposis coli	APC					
phosphomevalonate kinase	A POP	1			4D N	203527 s at 4
member	DANA	10654	NP 006547	NM 006556	4D Z	203515_s_at -
tumor necrosis factor receptor superfamily.	TNFRSFIB	7133	NP_001057	Nix1_001000		
EPH receptor A2	EPHA2					
PPAR binding protein	PPARBP	5469	NP_004765		AD IN	
translokin	PIG8	9702	NP_055494			
Sec61 gamma subunit	SEC61G	0	NP_055117		AD E	=
subcomplex unkn						203484 at
NADI Jak	NDI JECI		NP 002485		4D	203478_at
Solute carrier organic anion transporter family, m	IOGAP2		NP 006624		4D	
piecksuin	SLCOOR	٥	NP 009187	NM_007256	4D	203472_s_at
precesum	PIEK		NP 002655	NM_002664	4D	
eschildependent kinase (CDC2like) 10	PLEK		NP 002655	NM_002664	4D	
presentint 1 (Aizneimer disease 3)	CDKIO	8558	NP 003665	NM_003674	4D	
presenting 1 (Ald)	PSENI		NP 015557	NM_007318	4D	
spermiding/spermin blidding I	SAT		NP_002961	NM_002970	4D	203455 s at
I IM domain binding 1	LDBI	8861	NP_003884	NM_003893	4D	
PKD2 interactor, golgi and endoplasmic reticulum	PGEAI	25776	NP_056188	NM_0153/3	ŧ	
cadherin 2, type 1, Neadherin (neuronal)	CDH2	1000	NP_001783			
ribonuclease P/MRP 30kDa subunit	RPP30	10556	NP 006404			
endopeptida			NID COCKET		40	203436 at
membrane metalloendopentidase (neutral	MME	4311	NP_009218	NM_007287	4D	203435_s_at
family with sequence similarity 9	FAM8A1	51439	NP 057339	NM_016255		203420 at
CDS2 miles	CD53	963	NP 000551		4D	203416 at
n aisierin	DSCR2	8624	NP 003711		4D	203405_at
pregnancy specific beta [glycoprotein 3	TF 000	7018	NP 001054		4D	203400 s at
Nacetylg	PSG3	5671	NP 066296	NM_021016	4D	203399 x at
X1		_				

203564 at	8	NM 004629	NP_004620	2189	FANCG	Fanconi anemia, complementation group C
203574 at	ŧ	NM 003611	NP 003602	8481	OFDI	oralfacialdigital syndrome
203575 at	4) 6	NM 001806	NP 0035/3	4783	NFIL3	nuclear factor, interleukin 3 regulated
203579 s at	4b	NM 003983	1001007	1439	CSNK2A2	casein kinase 2, alpha prime polypeptide
203580 s at	4	NM 003083	NP 0039/4	9057	SLC7A6	solute carrier family 7 (cationic amino acid trans
203583 at	3 6	NIM 014044	NP 0039/4	9057	SLC7A6	solute carrier family 7 (cationic amino acid trans
203597 s at	4 6	NW 007197	NP 054763	25972	UNC50	unc50 homolog (C. elegans)
202500	ŧ	NM_00/18/	NP_009118	11193	WBP4	WW domain binding protein 4 (formin binding protei
202077_3_dt	ŧ	NM_007187	NP_009118	11193	WBP4	WW domain binding protein 4 (formin binding
20360 / at	40	NM_014937	NP_055752	22876	INPPSF	inositol polyphosphotosphosphotospho
203611_at	4D	NM_005652	NP_005643	7014	TERF2	telomeric report binding fortunates r
203614_at	4D	NM_021645	NP_067677	9724	UTP14C	UTP14, U3 small nucleolar ribonucleoprotein,
203616_at	4D	NM_002690	NP 002681	5423	POLB	nothing (DNA disease)
203617 x at	4D	NM_005229	NP_005220	2002	ELK1	ELK I, member of ETS oncome family.
203034 s at	40	NM_001876	NP_001867	1374	CPTIA	carnitine palmitov/transferase 1 A (line)
203635 at	40	NM_006052	NP_006043	10311	DSCR3	Down syndrome critical person game 3
203650 at	5	NM_006404	NP_006395	10544	PROCR	protein C receptor, endothelial (EPCR)
203033 S at	ŧ	NM_004645	NP_004636	8161	COIL	coilin
203035_at	ŧ	NM_006297	NP_006288	7515	XRCCI	Xray repair complementing defective repair in
203667_at	4D	NM_004607	NP_004598	6902	TBCA	tubulinspecific chaperone a
203668_at	Đ	NM_006715	NP_006706	4123	MAN2CI	mannosidose olpho oloco o
203672_x_at	45	NM_000367	NP_000358	7172	TPMT	thionurine Smethylrene ferroe
203675_at	4	NM_005013	NP_005004	4925	NUCB2	nucleobindin 2
203676 at	ŧ	NM_014967	NP_055782	22909	KIAA1018	KIAA1018 protein
203600 at	1 6	NM 006858	NP 006849	11018	TMED1	transmembrane emp24 domain containing 1
203790 at	ŧ	NM_006322	NP_006313	10426	TUBGCP3	tubulin, gamma complex associated protein 2
203708 - at	€	NM 017722	NP 060192	55621	FLJ20244	hypothetical protein FLJ20244
205/08_at	ŧ	NM_002600	NP_002591	5142	PDE4B	phosphodiesterase 4B, cAMPspecific
203737 s at	ð	NM 0148/8	NP 055693	9933	KIAA0020	KIAA0020
-out of at	+5	790CIO ININ	NP 055877	23082	PPRC1	norovicomo molificación

						gamma
203745_at	45	NM_005333	NP_005324	3052	HCCS	holocytochrome c synthase (cytochrome c
203748_x_at	4D	NM 016839	NP 058523	5037	BDMC	hemelyase)
			02022	7666	RBMSI	RNA binding motif, single stranded interacting
203753_at	4D	NM_003199	NP 003190	6925	TCF4	pio
203760 s at	4D	NM_006748	NP 006739	6503	SI.A	Spelibackerton
203761_at	4D	NM_006748	NP 006739	6503	A IS	Spoliton-donton
203765_at	4D	NM_012198	NP 036330	25801	GCA	orencedoir EEL
203766 s at	4D	NM_012134	NP 036266	25802	LMODI	grandin binding protein
203781_at	4D	NM_004891	NP 004882	9553	MRPL 33	mitochordist ib
203795_s_at	4D	NM_020993	NP 066273	605	BCI.7A	Boll CH /harb 23
203800_s_at	4D	NM_022100	NP 071383	63931	MRPS14	mitochandrid ribana /A
203802 x at	4D	NM_018044	NP_060514	55695	NSUNS	NOLL NOBS (Sim demail from S14
203810 at	4D	NM_007034	NP_008965	11080	DNAJB4	Dnal (Hendf) homolog subferrille in the control of
203813_s_at	40	NM_003062	NP_003053	6586	SLIT3	slit homolog 3 (Procentials)
203826 s at	45	7	NP_006538	10643	IMP3	IGFII mRNAbinding protein 3
203627 at	ŧ	NM_007371	NP_031397	8019	BRD3	bromodomain containing 3
203027_at	±	NM_017983	NP_060453	55062	WIPI49	WD40 repeat protein Interacting with
203831_at	4D	NM_014925	NP 055740	22864	KIAA1002	KIAAI002 protein
203832 at	4D	NM_003095	NP_003086	6636	SNRPF	small nuclear sibanual
203835 at	4D	NM_005512	NP_005503	2615	GARP	of ventration A renetition and the polypeptide I:
203840 at	4D	NM_003666	NP_003657	8548	BLZFI	basic lencine zimer miclosy factor 1 (15 x 1)
203855 at	4D	NM_014969	NP_055784	22911	WDR47	WD repeat domain 47
203857_s_at	45		NP_006801		PDIR	for protein disulfide isomeraserelated
203868 S at	40		NP_001069	7412	VCAMI	vascular cell adhesion molecule 1
2038/1 at	ŧ		NP_056485	26168	SENP3	SUMO1/sentrin/SMT3 specific protesse 3
-02000_at	ŧ	NM_005694	NP_005685	10063	COX17	COX17 homolog, cytochrome c oxidase assembly
203885 at	4D	NM_014999	NP_055814	23011	RAB21	RABOI member BAS
at		NM_001348	NP_001339		DAPK3	deathassociated protein kinase 3
203693_at	£	NM_016283	NP_057367	6880	TAF9	TAF9 RNA polymerase II, TATA box binding
203900_at	4D	NM_024547				pioceni

203904_x_at	4D	NM 002231	NP 002222	2722	11 A 11	T
203910 at	ŧ)	NM 004815	NP 004806	9411	PARCI	Kangai I (suppression of tumorigenicity 6, prostat
203913_s_at	4D	NM_000860	NP 000851	3248	HPGD	hydroxygratical and the state of the state o
203914 x at	4D	NM_000860	NP_000851	3248	HPGD	hydroxyprostaglandii dehydrogenase 15(NAD)
203919 at	4	NM_003195	NP_003186	6919	TCEA2	transcription elongation factor A (SII) 2
203921_at	ŧ	NM_004267	NP_004258	9435	CHST2	carbohydrate (Nacetylglucosamine6O) sulfotransfera
202026 a at	ŧ	NM_00108/	NP_001678	513	ATP5D	ATP synthase, H+ transporting, mitochondrial F co
203930_S_at	ŧ	NM_004994	NP_004985	4318	MMP9	matrix metalloproteinase 9 (gelatinase B, 92kDa
203938_s_at	ŧ	NM_005679	NP_005670	9013	TAFIC	TATA box binding protein (TBP)associated
203943 at	45	NM_004798	NP 004789	9371	KIF3B	kinesin family member 1B
203944 x at	\$	NM_007049	NP_008980	11120	BTN2A1	butyrophilin, subfamily 2 member 41
2020CA =	ŧ	NM_001299	NP_001290	1264	CNNI	calponin I, basic, smooth muscle
203966_s_at	4D	NM 021003	NP 066283	5494	NMI	Nmyc (and STAT) interactor
203968_s_at	4D	NM_001254	NP_001245	990	CDC6	magnesiumdep CDC6 cell division cycle 6 homolog (S.
203973_s_at	4D	NM 005195	NP 005186	1052	CERPO	CC A AT/
203978_at	4D	NM_002484	NP_002475	4682	NUBPI	nucleotide binding protein 1 (MinD homolog, E.
203981_s_at	4	NM_020323	NP_064719	5826	ABCD4	ATPbinding cassette, subfamily D (ALD), member 4
204000 at	ŧ	NM_016194	NP_057278	10681	GNB5	guanine nucleotide binding protein (G protein) be
204004 at	ŧ	NM_002583	NP_002574	5074	PAWR	PRKC, apoptosis, WT1, regulator
20401 at	ŧ	NM_005842	NP 005833	10253	SPRY2	sprouty homolog 2 (Drosonhila)
204020 at	45	NM_005859	NP_005850	5813	PURA	purinerich element binding protein A
204021_s_at	4D	NM_005859	NP_005850	5813	PURA	purinerich element hinding protein A
204042 at	ŧ	NM 006646	NP_006637	10810	WASF3	WAS protein family, member 3
204053_x_at	£	NM_000314	NP_000305	5728	PTEN	phosphatase and tensin homolog (mutated in multipl
	100	141C000 141VI	NP 000305	5728	PTEN	phosphatase and tensin homolog (mutated in

aldehyde dehydrogenase 6 family manhan A I	ALDH6A1	4329	NP_005580	NM_005589	15	204290 S at
pim2 oncogene	PIM2	11040	NP_006866	C/8900 IMN		
S100 calcium binding protein A2	S100A2		NP 003969	M 000076		
cyclindependent kinase 2	CDK2		NP 005050	NM 005078		
GULP, engulfment adaptor PTB domain containing I	GULPI	•	NF_03/399	NM 001708		
chromosome 7 open reading frame 23	C7orf23		NP 077291	NM 016315		a
solute carrier family 31 (copper transporters), me	SLC31A2		NP 001851	0001000 MIN	40	204215 at
BTB and CNC homology 1, basic leucine zipper trans	BACHI		NE OOLL	NIM DOING		204204 at
peptidylprolyl isomerase D (cyclophilin D)	Clidd	_	NP_005029	NM 001186	4) 5	204194 at
peptidylprolyl isomerase D (cyclophilin D)	DIID	5481	NP_005029	NM_005038		204186 S at
RNA binding motif protein 14	RBM14	10432	NP_006319	NM 006528		204195 v at
kinetochore associated 2	KNTC2	10403	NP_006092	NM_006101		204162 at
transcription factor Dn1	TFDPI	7027	NP_009042	NM_007111		204147 s at
enolase superfamily member 1	ENOSFI	55556	NP_059982	NM_017512	4D	204143 s at
enolase superfamily member 1	ENOSFI	55556	NP_059982	NM_017512		204142_at
tyrosylprotein sulfotranetarasa 1	TPSTI	8460	NP_003587	NM_003596	4D	204140_at
forkhead box O3A	FOXO3A	2309	NP_001446	NM_001455	4D	204132 s at
forkhead box O3A	FOXO3A	2309	NP_001446	NM_001455	4D	204131_s_at
replication factor C (activator 1) 3, 38kDa	RFC3	5983	NP_002906	NM_002915	4D	204128_s_at
renlication factor ((cational 1) 2 222	RFC3	5983	NP_002906	NM_002915	4D	204127_at
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex	NDUFAFI	51103	NP_057097	NM_010013	ŧ	an Carlo
prolyl endopeptidase	PREP	5550	NP_002717	NM 002726		204117 at
RNA binding motif protein. Xlinked?	RBMX2	51634	NP_057108	NM_016024		201117 0
RNA binding motif protein, Xlinked 2	RBMX2	51634	NP_057108	NIVI 010024		204008 of
solute carrier family 5 (sodiumdependent vitamin t	SLC5A6	8884	NP 066918	C60170 IMM		204097 s at
glycine, glutamate, thienylcyclohexylpineridinehin	KIAA0562	9731	NP 055519	MM 021002		204087 s at
topoisomerase I binding, arginine/serinerich	TOPORS	10210	NP 005793	709CO MIN	40	204075 s at
Meis I, myeloid ecotropic viral integration site I	MEIS1	4211	NP_002389	NM 005903		204071 s at
unc51like kinase 2 (C. elegans)	ULK2	9706	NP 055498	NM 014683	ŧ	204069 54
unc51like kinase 2 (C. elegans)	ULK2	9706	NP 055498	NM 014683	3 6	201063 6 04
munpi			100	NIM DIACOS	45	204062 s at

nuclear pore complex interacting protein	NPIP	9284	NP 008916	C86000 IAINI	40	-0.000 v at
breast cancer 1, early onset	BRCAI	2/0	MF_009220	NA 000000	Ì	204538 v ot
myosin VA (heavy polypeptide 12, myoxin)	MYOSA	4044		NM 00705	45	204531 s at
domain) domain family, member 8 (with GRAM	100		NID OOOSS	NM 000250	4D	204527 at
The cell division cycle 7 (S. cerevisiae)	TBCID	11138	NP 008994	NM 007063	4D	204526_s_at
CDC7 - II 4:	CDC7	8317	NP 003494	NM_003503	4D	204510_at
protein phosphores 2 (female agonts)	PPP3R1	5534	NP 000936	NM_000945	4D	204507_s_at
BH3 interacting domain double consist	BID	637	NP_001187	NM_001196	4D	204493 at
Transmembrane protein 15	TMEM15	22845	NP_055723	NM_014908	4D	204488_at
enolase 3 (beta muscle)	ENO3	2027	NP_001967	NM_001976	4D	204483_at
bromodomain and PHD finger containing	BRPF1	7862	NP 004625	NM_004634	Đ	204481_at
anolinoprotein Al	APOAI	335	NP_000030	NM_000039	4D	204450 x at
arachidonate clinoxumaning protein	ALOX5	240	NP_000689	NM_000698	4D	204445 s at
PH domain containing and	pp9099	80301	NP 079477	NM_025201	4D	204436_at
muskeim i, illuraceilular mediator containing kelc				NM_014778	4D	204435_at
aponpoproteiii CI	MKIN	4289	NP 037387	NM_013255	4D	204423_at
anolinoprotein CI	APOCI	341	NP_001636	NM_001645	4D	204416_x_at
neurofilament heavy nolvegatide 2001.5	NEFH	4744	NP_066554	NM_021076	4D	204412 s at
echinoderm microtubula accominate o	EML2	24139	NP_036287	NM_012155	4D	204398 s at
G proteincounled recentar binace 5	GRK5	2869	NP_005299	NM_005308	4D	204396 s at
mitochondrial ribosomal protein 62	MRP63	78988	NP_076931	NM_024026	4D	204387 x at
fibroblast growth factor recontor 2 /- L	FGFR3	2261	NP_000133	NM_000142	40	2043/9 s at
breast carcinoma amplified sociona-	BCASI	8537	NP_003648	NM_003657	40	204378 at
Varhinding protein	VprBP	9730	NP_055518	NM_014703	4D	204376 at
KHtype splicing regulatory protein (FUSE binding	KHSRP	8570	NP_003676	NM_003685	÷	2045/2_s_at
ATP/GTPbinding protein	HEAB	10978	NP_006822	NM_006831	40	2043/0 at
nohmanid	GTF3C2	2976	NP_001512	NM_001521	45	204366 s at
TNF receptorassociated factor 5	TRAF5	7188	NP 004610	NM_004619	40	204352 at
S100 calcium binding protein P	S100P	6286	NP_005971	NM_005980	4	204351 at
collagen, type XVI alpha	COL16A1	1307	NP_001847	NM_001856	4D	204345 at
zinc finger protein 202	ZNF202	7753	NP_003446	NM_003455	45	204327 s at
Banylate cyclase B				NM_002450	4D	204326_x_at
natrimetic pentide records D	NPR2	4882	NP 003986	NM_003995	40	204310 s at

HI histone family member v	HIFX	8971	NP 006017	NM_006026	£	204802 S at
hypothetical protein FLJ13639	FLJ13639	/9/58	NP 0/8981	ININ 024703	3	20/00/20 200
G proteincoupled receptor associated sorting prote	OPKASPI	7/3/	NI 02022	SOUTH OF THE	ה ה	204800 s at
Fas (TNF receptor superfamily, member 6)	FAS	0777	100000 Bi	NM 014710	4D	204793 at
ras (1 NF receptor supertamily, member 6)	125	255	NP Onna	NM 000043	4D	204781 s at
econopic vital integration site 2A	EAC	355	NP 000034	NM 000043	4D	204780 s_at
ecotronic viral integration disco	EVI2A	2123	NP 055025	NM_014210	4D	204774 at
transcription termination factor, RNA polymerase	TTFI	7270	NP_031370	NM_00/344	ŧ	204771_3_at
nudix (nucleoside diphosphate linked moiety X)type	NUDIT	1321	141 002445		à	204771
farnesyltransferase, CAAX box, beta	FNTB	2342	NP 002019	NM 002020	4 6	204766 s at
(prostaglandi				OCCCOO MIN	4	204764 at
prostaglandinendoperoxide synthase 2	PTGS2	5743	NP_000954	NM_000963	4D	204748_at
isoleucinetRNA synthetase	IARS	3376	NP_038203	NM_013417	40	204744 S at
chondroitin sulfate proteoglycan 4 (melanomaassoci	CSPG4	1464	NF_001888	Min_001037		
NCK adaptor protein 1	NCK1	4690	NP 006144	NIM 001007	4D	204736 s at
coagulation factor V (proaccelerin, labile factor)	F5	2153	NP 000121	NM 006152	4 6	204725 s at
hypothetical protein MGC29875	MGC29875	2/042	NF 033203	90CF10 1414	È	204713 s of
cadherin 5, type 2, VEcadherin (vascular epitheliu	CDHS	1003	NE_001700	882V10 WN	45	204700 x at
(3oxo5			ND OOLSO	NM 001705	45	204677 at
steroid5alphareductase, alpha polypentide 1	SRD5A1	6715	NP_001038	NM_001047	£	3E_C/0#02
ring finger protein 24	RNF24	11237	NP_009150	NM_007219	ŧ	204675 at
adenosine deaminase	ADA	100	NP_000013	NM 000022	ŧ	204668 at
golgi SNAP receptor complex member	GOSR1	9527	NP_004862	NM_004871	ŧ	304620 H
nuclear receptor subfamily 4, group A, member 2	NR4A2	4929	NP_006177	NM 006186	ŧ	18 X 770+07
adrenocortical dysplasia homolog (mouse)	ACD	65057	NP_075065	NM_022914	ŧ	204617 S at
dickkopf homolog 1 (Xenopus laevis)	DKKI	22943	NP_036374	NM_012242	40	204002 at
protein (peptidylprolyl cis/trans isomerase) NIMA	PIN4	5303	NP_006214	NM_006223	ŧ	204507 X at
KIAA0831	KIAA0831	22863	NP 055739	NM 014924	£	204590 at
FK506 binding protein 5	FKBP5	2289	NP 004108	NM 004117	ŧ	204569 at
protein phosphatase I, regulatory subjunit 3D	PPP1R3D	5509	NP_006233	NM_006242	40	204555 s at
protein phosphatase 1, regulatory subunit 313	PPP1R3D	5509	NP_006233	NM_006242	40	204554 at
HermanskyPudlak syndrome 5	HPS5	11234	NP_009147	NM_007216	4D	204544 at
eukaryotic translation elongation factor I alaka a	EEF1A2	1917	NP 001949	8C6100 IMM	t	-04040 at

APPENDIX I

	MASI	11157	NP 009011	NM 007080	4D	205036 at
cyclin E2	CCNE2	9134	NP_004693	NM_004702	40	200034 at
transforming growth factor, alpha	TGFA	7039	NP 003227	NM 003236	ŧ	205024 at
transforming growth factor, alpha	TGFA	7039	NP 003227	NM 003236	ŧ	05016 at
trefoil factor 1 (breast cancer, estrogeninducible	TFF1	7031	NP_003216	C77500 IAIN	à	205005 at
Nmyristoyltransferase 2	NMT2	9397	NP 004/99	900+00 ININ	A) đ	205000 at
Nmyristoyltransferase 2	NMT2	9397	NP 004/99	MIN 001000	40	205006 s at
cyclindependent kinase 5, regulatory subunit 1 (p3	CDK5R1	1088	NP 0038/6	000000 MINI	40 6	205005 s at
integrin, beta 4	ITGB4	3691	NP 000204	200200 MIN	à é	204006 6 91
clock homolog (mouse)	CLOCK	95/5	NP 004889	000000 MIN	É	204989 s at
neutrophil cytosolic factor 1 (47kDa, chronic gran	NCF1	4687	NP 000256	C97000 IAIN	ð 8	204980 at
myeloid cell nuclear differentiation antigen	MNDA	4332	NP 002423	INM 002452	ŧ	204061 6 94
pololike kinase 3 (Drosophila)	PLK3	1263	NP 004064	NW 004073	1	04050 at
origin recognition complex, subunit 5like (yeast)	ORC5L	5001	NP_002544	NW 00233	3 8	204059 04
intercellular adhesion molecule 3	ICAM3	3385	NP_002153	701700 IAIN	3 6	204957 at
protein tyrosine phosphatase, receptor type, G	PTPRG	5793	NP 002832	NM 002172	3 8	204949 at
inhibin, beta A (activin A, activin AB alpha polyp	INHBA	5024	100200	TYRCOO MIN	5	204944 at
tollike receptor 2	ILIN2	7097	NID 002102	COLCOO MIN	45	204926 at
Common	TIDO	7007	NP 003255	NM 003264	4D	204924 at
myeloid/lymphoid or mixedlineage leukemia	MLLT3	4300	NP_004520	NM_004529	ŧ	70491/_S_at
eukaryotic translation elongation factor ensilon	EEFIEI	9521	NP_004271	NM 004280	ŧ	204017
amine oxidase, copper containing 3 (vascular adhes	AOC3	8639	NF_003/25	MM_00373#	3 6	204005 6 04
HUS1 checkpoint homolog (S. pombe)	HUSI	3364	NP 004498	10C400/	è	204894 s at
Rho GTPase activating protein 25	ARHGAP25	9938	NP 053697	700410 MIN		204884 e at
immature colon carcinoma transcript	ICTI	3396	NP 001536	CHCLOO ININ	£ 6	204887 at
baculoviral IAP repeatcontaining 1	BIRCI	4671	NP_00452/	0000 MINI	100	204868 at
baculoviral IAP repeatcontaining 1	BIRCI	4671	NP 004527	000500 MINI	5 6	204861 s at
protein kinase, cAMPdependent, regulatory, type	PRKAR2A	5576	NP_004148	NM_004537	è	204860 s at
early endosome antigen 1, 162kD	EEA1	8411	NP_003557	NM 003566	4 6	204842 v at
processing of precursor 5, ribonucleuse P/MRP subu	POP5	51367	NP_05/002	816CIO MN	i	201000
mutL homolog 3 (E. coli)	MLH3	27030	NP_055196	NM 014381	à é	204839 at
major matocompanionney complex, class I. F.						W . V . V

homogentisate 1,2dioxygenase (homogentisate oxidas	HGD	3081	NP_000178	NM 00018 /	ŧ	1 1 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
polymerase (RNA) III (DNA directed) polypeptide F,	POLR3F	1 2				
apolipoprotein H (beta2glycoprotein 1)	APOH					l
serine/threonine kinase 17b (apoptosisinducing)	SIKI7B	-				205216 s ar
vrel reticuloendotheliosis viral oncogene homolog						205214 at
prosprioserine prospiratase				NM 006509	4D	205205 at
nhosphosering phosphotos	PSPH	5723	NP 004568	NM_004577	4D	205194_at
vmaf musculoaponeurotic fibrosarcoma oncogene	MAFF	23764	NP_036455	NWI_012323		1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
retinitis pigmentosa 2 (Xlinked recessive)	RP2					205193 at
a disintegrin and metalloproteinase domain 8	ADAM8	101				205101 0+
CD58 antigen, (lymphocyte functionassociated antig	CD58	965				205170
leucine rich repeat neuronal 5	LRRN5	10446	NP_006329			205173 v at
chloride channel 4	CLCN4	1183	NP_001821			205150 04
hypothetical protein FLJ22269	FLJ22269	84179				205149 s at
uronyl2sulfotransferase	UST	10090	NP_005706			205144 ot
heat shock 10kDa protein 1 (chaperonin 10)	HSPE1		NP_002148			205139 s at
sai cogiycan, beta (45KDa dystrophimassociated glyc						205122
romy behave receptor i	SCCB		NP 000223	NM_000232	4D	205121_at
formul postide	FPR		NP 002020	NM_002029	4D	205119 s_at
syntanhilin	SNPH		NP 055538	NM_014723	4D	205104_at
solute carrier family 26 (culfety to	SLC26A2	1836	NP_000103	NM_000112	45	205097_at
Nacetylglucosamine I phosphodiester alphaNacetylgluc	NAGPA	51172	NP_05/540	MM_010200		
interleukin I, beta	ILIB		NP 00056/	95C910 WIN		205090 s at
AT rich interactive domain 4A (RBP1like)	ARID4A	5926	NP 002883	769700 IAIN		205067 at
primase, polypeptide 1, 49kDa	PRIMI		NP 00093/	C00C00 MIN		205062 x at
oncogen			NID OOOOOO	NM 000046	4D	205053 at
vkit HardyZuckerman 4 feline sarcoma viral	KIT	3815	NP_000213	NM_000222	4D	205051 s_at
				NM 003832		20-010 S at
glucosamine (UDPNacetyl)2epimerase/Nacetylmannosam	GNE	10020	TAT_005407			205010
(S.			NID 005467	NM 005476	45	205042 at
		-	-			

2000						
205222_at	ŧ	NM_001966	NP_001957	1962	ЕННАДН	enoylCoenzyme A, hydratase/3hydroxyacyl Coenzyme A
205227_at	4D	NM_002182	NP_002173	3556	ILIRAP	interleukin 1 receptor accessory protein
205248 at	4D	NM_005128	NP_005119	9980	C21orf5	chromosome 21 open reading frame 5
205279 s at	40	NM_000824	NP_000815	2743	GLRB	glycine receptor, beta
205285 s_at	4D	NM_001465	NP 001456	2533	FYB	FYN binding protein (FYB120/130)
205286_at	4D	NM_003222	NP_003213	7022	TFAP2C	transcription factor AP2 gamma (activating enhance
205296_at	4D	NM_183404	NP_899662	5933	RBL1	retinoblastomalike 1 (p107)
205310_at	4D	NM_012066				4
205317_s_at	4D	NM_021082	NP_066568	6565	SLC15A2	solute carrier family 15 (H±/pentide transporter)
205318_at	4D	NM_004984	NP_004975	3798	KIF5A	kinesin family member 5A
205327_s_at	4D	NM_001616	NP_001607	92	ACVR2	activin A receptor, type []
205329_s_at	4D	NM_003794	S82500 dN	8723	SNX4	sorting nexin 4
205340_at	4D	NM_014797	NP_055612	9841	ZBTB24	zinc finger and BTB domain containing 24
205346_at	4D	NM_006927	NP_008858	6483	ST3GAL2	ST3 betagalactoside alpha2,3sialyltransferase 2
205350 at	40		NP_068832	11013	TMSNB	thymosin, beta, identified in neuroblastoma cells
202229 at	ŧ		NP 000817	2891	GRIA2	glutamate receptor, ionotropic, AMPA 2
2053/1 S at	100		NP_001909	1629	DBT	dihydrolípoamide branched chain transacylase E2
2055/4 at	ŧ		NP_003054	6588	SLN	sarcolipin
205384_at	ŧ	NM_005031	NP_005022	5348	FXYDI	FXYD domain containing ion transport regulator
205403_at	4D	NM_004633	NP_004624	7850	ILIR2	interleukin I receptor, type II
205406 s at	4D	NM_017425	NP_059121	53340	SPA17	sperm autoantigenic protein 17
205414 s at	4D	NM_014859	NP_055674	9912	K1AA0672	KIAA0672 gene product
205422_s_at	4D	NM_004791	NP_004782	9358	ITGBLI	integrin, betalike 1 (with EGFlike repeat domains)
205423_at	4		NP_001118	162	APIB1	adaptorrelated protein complex 1, beta 1 subunit
205433 at	4	NM_000055	NP_000046	590	BCHE	butyrylcholinesterase
205436 s at	40			3014	H2AFX	H2A histone family, member X
205438_at	45		NP_008970	11099	PTPN21	protein tyrosine phosphatase, nonreceptor type 21
205441 at	4		NP_078854	79629	FLJ22709	hypothetical protein FLJ22709
205481 at	100			8	MFAP3L	microfibrillarassociated protein 3like
205 182 at	16				ADORAI	adenosine A1 receptor
200462 X at	#	NM_013306	NP_037438	29907	SNX15	sorting nexin 15

coagulation factor VIII, procoagulant component	F8	2157	NP_000123	NM_000132	40	205/56_s_at
biphenyl hydrolaselike (serine hydrolase; breast e	BPHL	670	NP_004323	NM_004332	ŧ	205/50 at
oncostatin M receptor	OSMR	9180	NP_003990	NM_003999	4	205/29 at
ciliary neurotrophic factor receptor	CNTFR	1271	NP_001833	NM_001842	40	205725_at
proopiomelanocortin (adrenocorticotropin/ betalipo	POMC	5443	NP_000930	NM_000939	5 6	205720_at
bone marrow stromal cell antigen 1	BSTI	683	NP_004325	NM_004334	4	205715 at
interleukin 17 receptor	IL17R	23765	NP_055154	NM_014339	4	205/07 at
mitogenactivated protein kinase kinase 6	MAP2K6	5608	NP_002749	NM_002758	45	205698 s at
maternal G10 transcript	G10	8896	NP_003901	NM_003910	45	205690 s at
BCL2related protein A1	BCL2A1	597	NP_004040	NM_004049	4D	205681_at
major histocompatibility complex, class II. DO bet	HLADOB	3112	NP_002111	NM_002120	45	205671 s at
complement component 4 binding protein alpha	C4BPA	722	NP_000706	NM_000715	4D	205654_at
small nuclear ribonucleoprotein polypentide G	SNRPG	6637	NP_003087	NM_003096	4D	205644 s at
centrosomal protein 1	CEPI	11064	NP_008949	NM_007018	4	205642 at
cytidine deaminase	CDA	978	NP_001776	NM_001785	4D	205627 at
multimerin 1	MMRNI	22915	NP_031377	NM_007351	40	205612_at
homeo box B5	нохв5	3215	NP_002138	NM_002147	40	200600 x at
TRAF interacting protein	TRIP	10293	NP_005870	NM_005879	4D	205598 at
FGFR1 oncogene partner	FGFRIOP	11116	NP_008976	NM_007045	8	205588 s at
lysosomalassociated membrane protein 3	LAMP3	27074	NP_055213	NM_014398	45	205569_at
carbohydrate (keratan sulfate Gal6) sulfotransfera	CHST1	8534	NP_003645	NM_003654	4D	205567_at
frataxin	FXN	2395	NP_000135	NM_000144	4D	205565 s_at
ribonuclease P/MRP 38kDa subunit	RPP38	10557	NP_006405	NM_006414	4D	205562_at
TNF receptorassociated factor 6	TRAF6	7189	NP_004611	NM_004620	4D	205558_at
deoxyribonuclease Hike 3	DNASEIL3	1776	NP_004935	NM_004944	4D	205554_s_at
G1 to S phase transition 2	GSPT2	23708	NP_060564	NM_018094	4D	205541 s at
cadherin 6, type 2, Kcadherin (fetal kidney)	CDH6	1004	NP_004923	NM_004932	4D	205532 s at
hyaluronan and proteoglycan link protein 1	HAPLNI	1404	NP 001875	NM_001884	40	205523_at
transcobalamin I (vitamin B12 binding protein, R	TCNI	6947	NP_001053	NM_001062	4	200015_at
villin I	VILI	7429	NP 009058	NM 007127	ŧ	205506 at
KIAA0408	NIAA0408	9119	INT OCCUPA	101110	10	

calcotic D (SEI D	6403	NP 002996	ZM 003005	£	206049 at
nucleolar protein 4	NOL4	8715	NP 003778	NM_003/8/	ŧ	206040 S at
RAB33A, member RAS oncogene family	RAB33A	9363	NP_004/85	100 DOLDO	4 6	206045 8 24
aspartoacylase (aminoacylase 2, Canavan disease)	ASPA	443	NP 000040	NM 000049	à t	206039 at
				NM 014645	ŧ	200000 at
ELK4, ETSdomain protein (SRF accessory protein 1)	ELK4	2005	NP_001964	NM_001973	ŧ	203994_BI
Tbox 2	ТВХ2	6909	NP_005985	NM_005994	ŧ	205004 S at
CDIC antigen, c polypeptide	CDIC	911	NP_001756	NM_001765	45	205002 205002
klotho	KL	9365	NP_004786	NM_004795	40	205067 at
histone 1, H4c	HIST1H4C	8364	NP_003533	NM_003542	40	205967 at
basic leucine zipper transcription factor ATFlike	BATF	10538	NP_006390	NM_006399	5	205965_at
DnaJ (Hsp40) homolog, subfamily A, member 3	DNAJA3	9093	NP_005138	NM_005147	4D	205963 s at
pyruvate dehydrogenase kinase isoenzeme 4	PDK4	5166	NP_002603	NM_002612	4D	205960_at
plexin B3	PLXNB3	5365	NP 005384	NM_005393	4D	205957 at
carbonic anhydrase [CA1	759	NP_001729	NM_001738	4D	205950 s at
vanin 2	VNN2	8875	NP_004656	NM_004665	4D	205922_at
solute carrier family 6 (neurotransmitter transpor	SLC6A6	6533	NP_003034	NM_003043	40	205921_s_at
solute carrier family 6 (neurotransmitter transpor	SLC6A6	6533	NP_003034	NM_003043	ŧ	205920 at
prepronociceptin	PNOC	5368	NP_006219	NM_006228	40	205901 at
Jak and microtubule interacting protein 2	KIAA0555	9832	NP_055605	NM_014790	#	20588 s at
leukemia inhibitory factor recentor	LIFR	3977	NP_002301	NM 002310	40	2058/6_at
AT rich interactive domain 3A (BRIGHT like)	ARID3A	1820	NP_005215	NM_005224	40	205865_at
solute carrier family 7 (cationic amino acid trans	SLC7A4	6545	NP_004164	NM_004173	40	205864 at
solute carrier family 18 (vesicular monoamine), me	SLC18A2	6571	NP_003045	NM_003034	É	200007_at
polymerase (DNA directed), gamma 2, accessory subu	POLG2	11232	NP_009146	NM 002027	ŧ	205011_at
tuftelin 1	TUFT1	7286	NP 064512	NM 020127	ŧ	205007 S at
TRAF3interacting Jun Nterminal kinase (JNK)activat	T3JAM	80342	NP_0/9504	NM 020228	ŧ	205007 5 4
tyrosine kinase, nonreceptor, 1	TNKI	8711	NP 003976	CSCCO INN	ð	205804 s at
receptor (calcitonin) activity modifying protein 2	RAMP2	10266	NP 003845	PCOCOO MINI	ð 8	205793 v at
coagulation factor XII (Hageman factor)	F12	2161	NP 000496	NM 000505	ð	205779 at
DEAD (AspGluAlaAsp) box polypeptide 18	DDAIG	0000			5	705774

chemokine (CXC motif) ligand 6 (granulocyte chemot	CACLO	00/2	-002307			
ryanodine receptor 3						206336 at
mase kinase kinase kinase kinase kina				NM 001036	4D	206306 at
mitogenactivated protein binace linear	MAP4K1	11184	NP 009112	NM_007181	4D	206296 x at
teratocarcinomaderived arough factor 1	TDGF1	6997	NP_003203	NM_003212		200280 S at
arginine vasopressin receptor IA	AVPRIA	552	NP_000697			200200 X at
YYI associated factor 2	YAF2	38				206250 5 01
G proteincoupled receptor 4	GPR4					206238 6 04
paired box gene 2	PAX2					206236 at
member					45	206229 x at
tumor necrosis factor receptor superfamily,	TNFRSF10C	8794	NP_003832	NM_003841	4	200222 at
carbonic anhydrase IV	CA4	762	NP_000708			200200
carbonic anhydrase IV	CA4					206200
Mphase phosphoprotein 9	MPHOSPH9	98	NP 073619			206200
unc5 homolog C (C. elegans)	UNC5C	8633				206205 at
crystallin, beta B1	CRYBBI	1414	NP_001878			206180 at
arginase, liver	ARGI	383	NP_000036			201205 at
adrenergic, beta2, receptor, surface	ADRB2					206177 6 4
pentaxinrelated gene, rapidly induced by IL1 beta	PTX3	6	NP 002843			206170 at
LIM homeobox 2	LHX2		NP 004780			206157 at
frizzled homolog 5 (Drosophila)	FZD5		NP_003459			206140 at
arylsulfatase B	ARSB		NP 000037			206136 at
early growth response 3	EGR3	1960	NP_004421			206129 s at
EPH receptor A4	EPHA4	2043	NP 004429			206115 at
histone 1, H3h	HIST1H3H	8357	NP 003527		45	206114 at
rasrelated C3 botulinum toxin substrate 3 (rho fam	RAC3	1880	NP 005045			206110 at
solute carrier family 22 (organic cation transport	SLC22A1LS	2003	NP_009036			206103 at
NELlike I (chicken)	NELLI	4/43	ND 000036	NM 007105		206097 at
high mobility group AThook 1	INUMI	1715	ND 006149	NM 006157		206089 at
urocortin	OCN	0315	CC1500 div	NM 002131		206074 s at
NAD31 Homolog C (S. cerevisiae)	I Compare	73/0	NP 003344	NM 003353	4D	206072 at
BADELL Trooide technolett bolybebilde A.	RADSIC	5889	NP 002867	NM 002876	4D	206066 s at
small nuclear ribania	SNRPAI	6627	NP 003081	NM_003090	4D	206055 s at
zinc finger protein \$10	ZNF510	22869	NP 055745	NM_014930	4D	206053 at
antig						

hypothetical protein FLJ20344	FLJ20344	55634	NP_060246	NM_017776	40	200085 at
NK2 transcription factor related, locus 5 (Drosoph	NKX25	1482	NP_004378	NM 004587	45	206582 -1
mitogenactivated protein kinase kinase kinase kina	MAP4K4	9448	NP 004825	NM 004834	3 8	206578 of
SMA3	SMA3	10571	NP_006771	NM 006/80	ŧ	206571 A at
nucleoporin 155kDa	NUP155	9631	NP 004289	NM 004298	ŧ	206565 2 64
SWI/SNF related, matrix associated, actin dependen	SMARCBI	6598	NP_003064	NIM 004208	5	206550 s at
4aminobutyrate aminotransferase	ABAT	18	NP 000654	NM 000663	ŧ	206532 at
maltaseglucoamylase (alphaglucosidase)	MGAM	8972	NP_004659	NM 004668	ŧ	22002Z at
sine oculis homeobox homolog 2 (Drosophila)	SIX2	10736	NP 058628	NM_016932	ŧ	206577 of
oculocutaneous albinism II (pinkeye dilution homol	OCA2	4948	NP_000266	C/7000_IMN	ŧ	200420 at
MBD2 (methylCpGbinding protein)interacting zinc fi	MIZF	25988	NP_056332	NM_OIDST/	ŧ	206495 S. at
L1M domain binding 2	LDB2	9079	NP_001281	NM 001290	ŧ	206405 at
transducinlike enhancer of split 3 (E(spl)) homologic	TLE3	7090	NP_005069	NM_005078	45	206472 s at
neurotrophic tyrosine kinase, recentor type 3	NTRK3	4916	NP_002521	NM_002530	4D	206462_s_at
rhodopsin (opsin 2. rod pigment) (retinitis migman	RHO	6010	NP_000530	NM_000539	4D	206454 s at
lin7 homolog A (C. elegans)	LIN7A	8825	NP_004655	NM_004664	4D	206440 at
coagulation factor II (thrombin) recentorlike I	F2RL1	2150	NP_005233	NM_005242	4D	206429 at
melanA	MLANA	2315	NP_005502	NM_005511	40	206427 s at
cyclic nucleotide gated channel alpha 1	CNGAI	1259	NP_000078	NM_000087	₽ D	206417_at
development and differentiation enhancing factor	DDEF2	8853	NP_003878	NM_003887	ŧ	200717 3 41
ubiquitin specific protease 6 (Tre2 oncogene)	USP6	9098	NP_004496	NM 004505	ŧ	206414 x at
CD19 antigen	CD19	930	NP_001761	NM_001770	4	206398 s at
retinoic acid receptor responder (tazarotene induc	RARRESI	5918	NP_002879	NM_002888	4D	206392 s at
phosphodiesterase 3A, cGMPinhibited	PDE3A	5139	NP_000912	NM_000921	40	206389 s at
serine (or cysteine) proteinase inhibitor, clade A	SERPINA7	6906	NP 000345	NM_000354	ŧ	206386 at
brainderived neurotrophic factor	BDNF	627	NP_001700	NM_001709	4	206382 S at
myogenic factor 6 (herculin)	MYF6	4618	NP 002460	NM_002469	£	206392 at
chemokine (C motif) ligand 1	XCLI	6375	NP_002986	NM_002995	4D	206365 at
suppressor of cytokine signaling 3	SOCS3	9021	NP_003946	NM_003955	4D	206359 at
amyloid P component, serum	APCS	325	NP_001630	NM_001639	40	206350 at
iduronate 2sulfatase (Hunter syndrome)	IDS	3423	NP_006114	NM_006123	4	206342 x at

solute carrier family A godina bissal	SLC4AI0	57282	NP_071341	NM_022058	4D	206830 at
zinc finger protein 430	ZNF430	80264	NP 079465	681C70 IMM	ŧ	206820 Y at
peripheral myelin protein 2	PMP2	33/3	NF 002000	1,0200 talvi	à	00830
sperm associated antigen 8	SPAG8	26206	NP 036568	NM 012436	ð	206826 at
				8C/C00 IMM	ŧ	200000 3 41
prodynorphin	PDYN	5173	NP_077722	NM 024411	ŧ	200000 at
WNT1 inducible signaling pathway protein 1	WISPI	8840	NP_003873	NM 003882	ŧ	206903 at
verba erythroblastic leukemia viral oncogene homol	ERBB4	2066	NP_005226	NM_005235	ŧ	200774_01
killer cell lectinlike receptor subfamily C, membe	KLRC2	3822	NP 002251	NM_002260	ŧ	206794 at
acrosomal vesicle protein 1	ACRV1	56	NP 001603	Z19100 IMM	ŧ	206705 at
thymosin, beta 4, Ylinked	TMSB4Y	9087	NP 004193	NM_004202	18	206/09 at
potassium voltagegated channel, shakerrelated subf	KCNA5	3741	NP_002225	NM_002234	ŧ	200702_at
sperm associated antigen 9	SPAG9	9043	NP_003962	NM_003971	6	206763 s at
homeo box C5	HOXC5	3222	NP_061826	NM_018953	40	206/39 at
cholinergic receptor, nicotinic, alpha polynemide	CHRNA4	1137	NP_000735	NM_000744	40	206735 at
jerky homologlike (mouse)	JRKL	8690	NP_003763	NM_003772	4D	206734 at
growth hormone regulated TBC protein 1	GRTP1	79774	NP_078995	NM_024719	4D	206712 at
erythrocyte membrane protein hand 4 1132-3	EPB41L3	23136	NP_036439	NM_012307	4D	206710_s_at
chromosome 6 onen reading frame 32	C6orf32	9750	NP_056948	NM_015864	4D	206707 x_at
endothelin receptor type B	EDNRB	1910	NP 003982	NM_003991	4D	206701 x at
HIVI Tat interacting protein 60kDa	HTATIP	10524	NP_006379	NM_006388	4D	206689 x at
cleavage and polyadenylation specific factor 4 20	CPSF4	10898	NP_006684	NM_006693	4D	206688 s at
o open semme name o				NM 025024	4D	206659_at
chromosome 20 open reading frame 2	C20orf3	57136	NP_065392	NM_020531	4D	206656_s_at
-5 -5 -5 Francis (Secondary) (Second 21)				NM_016384	4D	206652_at
Shydroxytryntamine (serotonin) recentor 70	HTR2B	3357	NP 000858	NM_000867	4D	206638 at
apolipoprotein B mRNA editing enzyme, catalytic	APOBEC3B	9582	NP_004891	NM_004900	4	200652_s_at
ADAMTSlike 2	ADAMTSL2	9719	NP 055509	NM_014694	40	206629 at
interleukin 18 receptor 1	IL18R1	8809	NP_003846	NM_003855	40	206618 at
ovolike I(Drosophila)	OVOLI	5017	NP 004552	NM_004561	4D	206604_at
lymphocyte antigen 96	LY90	C+0C2	141 000 IV			200

glutamate receptor, ionotropic, Nmethyl	GRIN2D	2906	NP 000827	NM_000836	4D	20/036 x at
OF				NM_006316		
zona pellucida binding protein	ZPBP	11055	NP_008940	NM_007009		
RAB27B, member RAS oncogene family	RAB27B	5874	NP_004154	NM_004163		at
gammaaminobutyric acid (GABA) A receptor, beta 1	GABRB1	2560	NF_000803	NM 000817		
protein predicted by clone 23882	HSU79303	29903	NP 037433	NM 000613		207010 at
delta sleep inducing peptide, immunoreactor	DSIPI	1831	NP 004080	100000 MIN		1
Scavenger receptor class F, member 1	SCARFI	8578	NP 003684	NM 003693		207001 v at
chemokine (CC motif) receptor 6	CCR6	1235	NP 004358	NM 003/63		
protein tyrosine phosphatase, receptor type, f pol	PPFIA2	8499	NP 003616	NM 003625		
contactin 2 (axonal)	CNTN2	6900	NP 005067	NM 005076		
glucose6phosphatase, catalytic (glycogen storage d	G6PC	2538	NP_000142	MM_000121		200070
RUN and SH3 domain containing 1	RUSCI	23623	NP 055143	NM 000151	3 5	206057 of
promelaninconcentrating hormone	PMCH	5367	NP_002665	NM 0026/4		206042 S at
sema domain, immunoglobulin domain (Ig), short bas	SEMAJE	9723	141 000000			206042
protein kinase C, alpha	PRKCA	07/0	ND 026562	NM 012431		206941 x at
GLEI RNA export mediatorlike (yeast)	OLEIL	2733	001490	NM 002737		206923 at
classI MHCrestricted T cell associated molecule	CKIAM	2722	NID 001400	NM 001400		206920 s at
tripartite motificontaining 25	LKIM25	00//	ND 063550	NM 019604		206914 at
sallike I (Drosophila)	SALLI	0299	NE 005073	NM 005020		206911 at
MAX dimerization protein 1	MXDI	4084	NF 002050	NM 002000		206893 at
STE20like kinase (yeast)	SLK	9/48	NI 000016	NM 007357		206877 at
cadherin 4, type 1, Rcadherin (retinal)	CDH4	0749	ND 05535	NM 014720		206875 s at
harakiri, BCL2 interacting protein (contains only	HRK	8739	NP 003/9/	NM 003000	45 6	206866 at
hypothetical protein FLJ20323	FLJ20323	54468	NP 0618/8	COOCTO TANK		206864 s at
homeo box A7	HOXA7	3204	NP 00882/	NM 010005		206860 s at
fructose1,6bisphosphatase 2	FBP2	8789	NP 003828	VICECOO IAIN	3 8	206848 at
phosphodiesterase 6H, cGMPspecific, cone, gamma	PDE6H	5149	NP_006196	NM_006205	ŧ	200041_at
afamin	AFM	173	NP_001124	NM 001133	ŧ	206941 04
sema domain, immunoglobulin domain (1g), short bas	SEMA3F	6405	NP_004177	NM_004186	ŧ	200632_S_at
				har online	Cirl	027 0 24

207421_at	2	207414 c at	207307 at	207380 at	207383 6 94	207364 at	207224	207325 x at	207307 at	207286 at	2072/9_s_at	207254 at	207254 S at	207252 at 162702	207251 01	207223 at	207225 21	207206 8 24	207205 at	207186 s at	207167 at	207158_at	207152_at	207122_x_at	201071_3_4	207020 at	20700C ot	207072 at	207064 s at	207055 at	207041_at	
ŧ	3 5	ð) f	3 5	45	ð	Ē	4) 6	15	ŧ	40	40	1	4	ŧ	ŧ	đ	3 6	45	ê ê	40	4D	4D	4D	ŧ	ð	ð	3	40	4	4D	
NM_001739	0/C200 INN	CZCOO INN	C/1000 INN	106500 MINI	130500 MIN	NM 001587	100000	880700 MN	711 0000C	NM_025009	NM_016365	NM_005073	NM_016936		NM 014048	880100 IMN	1400007	NM DOLOIS	NIM 001917	NM 004459	85C/00 WN	NM_001644	NM 006180	NM_001054	NIN_000633	NM 005835	CCOCOO IAIN	120000 MIN	NM 009500	NM 004767	NM_00610	
NP_001730		NP 000514	NP 000164	NP 003952	NP 0015//	NP 003233	M _004777	NP 000839	NP_004133	NP_079285		NP_005064	NP_058632	NP_005916	NP_055463	NP_001079	NP_00088	NF_001808	ND 001000	ND 004249	NID OOLOGO	NP 001635	NP 006171	NP_001045	NP_005826	NP_006503	NP_005844	NE 027/20	NID 022700	ND DOOR	NP 006601	
763	5046	3239	2811	9028	1527	7048	100	3338	4328	9662		6564	29855	4225	9666	15	239	1089	2100	8666		339	4915	6799	10246	6291	8807	514	7203	2000	10747	
CA5A	PCSK6	HOXD13	GPIBA	RHBDL1	CXort2	TGFBR2	MAGEAI	HTR2C	MMPLI	KIAA0635		SLC15A1	UBNI	MEPIB	DZIP3	AANAT	ALOX12	CEACAM4	FALZ	IGSF2		APORECI	NTRK2	SULT1A2	SLC17A2	SAA4	IL18RAP	AUC2	GPK3/LI	INITAL Z	MASPO	
carbonic anhydrase VA, mitochondrial	proprotein convertase subtilisin/kexin type 6	homeo box D13	glycoprotein lb (platelet), alpha polypeptide	rhomboid, veinletlike I (Drosophila)	chromosome X open reading frame 2	transforming growth factor, beta receptor II (70/8	metanoma antigen family A, 1 (directs expression o	Shydroxytryptamine (serotonin) receptor 2C	matrix metalloproteinaselike 1	KIAA0635		solute carrier family 15 (o)igopeptide transporter	ubinuclein I	meprin A, beta	zinc finger DAZ interacting protein 3	arylalkylamine Nacetyltransferase	arachidonate 12lipoxygenase	carcinoembryonic antigenrelated cell adhesion mole	tetal Alzheimer antigen	immunoglobulin superfamily, member 2	po	anolinoprotein B mBNA addition	neurotrophic torogina kings of the Los, puctionples	sulfotransferase family cytosolic 14 phonologic	solute carrier family 17 (sodium phosphate), membe	serum amyloid A4, constitutive	interleukin 18 receptor accessory protein	amine oxidase, copper containing 2 (retinaspecific	Oprotein coupled receptor 37 like 1	mannanoinding tectin serine protease 2	mananhindin la di	Desparate

	KRTHR3 keratin hair hacia 3	3889	NP 002273	NM_002282	40	207069 at
mitogenactivated protein kinase kinase 3	MAP2K3	5606	NP 002747	95/700 IMN	ŧ	
a disintegrin and metalloproteinase domain 21	ADAM21		NP 003804	NM 003813	3 8	207667 s at
Tbox 1	TBXI	6899	NP 005983	NM 005992	ŧ	207665 at
SH3 multiple domains I	SH3MD1	9644	NP_055446	NM 014631	ŧ	207662 S at
myelinassociated oligodendrocyte basic protein	MOBP	4336	NP 006492	10C000 MN	ŧ	207661
transportin l	TNPOI	3842	NP_002261	NM 002270	ŧ	207650 S at
Bcell linker	BLNK	ľ	NP_037446	NM_013314	ŧ	207657 3 41
keratin, hair, acidic, 7	KRTHA7	8688	NP 003/61	1000 100 ININI	3 8	207655 0 01
tumor necrosis factor receptor superfamily, member	TNFRSFIA	7132	NP_001056	NM 001083	ŧ	207640 at
protease, serine, 7 (enterokinase)	PRSS7		NP_002763	NM_002772	ŧ	207643 G CT
serine (or cysteine) proteinase inhibitor, clade 1	SERPINI2	5276	NP_006208	NM 006217	ŧ	207628 at
corebinding factor, runt domain, alpha subunit 2:	CBFA2T2	9139	NP_005084	NM_005093	ŧ	207625 s at
hypocretin (orexin) receptor	HCRTRI	3061	NP_001516	NM_001525	40	20/019 at
calcium/calmodulindependent protein kinase (CaMki	CAMK2A	815	NP_057065	NM_015981	ŧ	207013_8_41
sulfotransferase family, cytosolic, 1B, member 1	SULTIBI	27284	NP_055280	NM_014465	ŧ	207613 at
chromosome I open reading frame 38	Clorf38	9473	NP_004839	NM_004848	45	20/5/1 x at
zinc finger protein 261	ZNF261	9203	NP_005087	NM_005096	40	207539 S at
membrane cofactor protein (CD46, trophoblastlympho	MCP	4179	NP_002380	NM_002389	ŧ	207547_X_at
numb homolog (Drosophila)	NUMB	8650	NP_003735	NM_003744	ŧ	207540 s at
exosome component 10	EXOSC10	5394	NP_002676	NM_002685	ŧ	207545 at
defensin, alpha 5. Paneth cellspecific	DEFA5	1670	NP_066290	NM_021010	ŧ	207549_at
bradykinin receptor B	BDKRB1	623	NP_000701	NM_000710	4	20/510_at
TBPinteracting protein	TIP120A	55832	NP_060918	NM_018448	4D	207483 s at
motilin	MLN	4295	NP_002409	NM_002418	4D	207473 at
DKFZp566H082 hypothetical protein DKFZp566H0824	DKFZp566H082	54744	NP_060005	NM_017535	40	20/4/0_at
galanin	GAL	51083	NP_057057	NM_015973	45	207466 at
glycophorin B (includes Ss blood group)	GYPB	2994	NP_002091	NM_002100	4D	207459 x at
chemokine (CC motif) receptor 9	CCR9	10803	NP_006632	NM_006641	40	207445 s at
solute carrier family 35 (1) Decalactors transport	SLC35A2	7355	NP_005651	NM_005660	4D	207439 s_at
solute carrier family 22 (organic cation transport	SLC22A2	2000	141 00000			

cytochrome P450, family 2, subfamily F, polypeptid	CIFZFI	1712	_000105			
		1673	ND 000765	NM 000774	4D	207913 at
THE PARTY OF THE				NM 004940	4D	207888 at
nuclear VCPlike	NVL	4931	NP 002524	NM 002533	4D	207877 s at
seizure related 6 homolog (mouse)like	SEZ6L	23544	NP_066938	NM_021115	4D	207873 x at
leukocyte immunoglobulinlike receptor, subfamily A	LILRA2	11027	NP_006857	NKI_006866	ŧ	20,007 at
chemokine (CXC motif) ligand 3	CXCL3	2921	NP_002081	NM 002090	ŧ	207057 at
interleukin 13	IL13	3596	NP_002179	NM 002188	ŧ	207044 81
cytochrome b5	CYB5	1528	NP 001905	NM 001914	ŧ	207844 x at
fibulin l	FBLN1	2192	NP_006478	NM 006487	ŧ	207835 at
BCL2/adenovirus E1B 19kDa interacting protein 1	BNIPI	662	NP_053581	NM_013978	8	207829_s_at
inhibitor of DNA binding 3, dominant negative heli	ID3	3399	NP_002158	NM_002167	ŧ	20/020_S_81
defensin, alpha 6, Paneth cellspecific	DEFA6	1671	NP_001917	NM_001926	ŧ	207814 at
ferredoxin reductase	FDXR	2232	NP_004101	NM_004110	40	20/815 s at
keratin 12 (Meesmann corneal dystrophy)	KRT12	3859	NP_000214	NM_000223	ŧ	207811 at
ataxin 2like	ATXN2L	11273	NP_059867	NM 017492	ŧ	207/98 s at
RAB1A, member RAS oncogene family	RAB1A	5861	NP_004152	NM_004161	ŧ	207791 s_at
keratin, hair, acidic, 3B	KRTHA3B	3884	NP_002270	NM_002279	ŧ	18 /8//02
arylsulfatase D	ARSD	414	NP_033667	NM_009589	ŧ	207767 at
presenilin I (Alzheimer disease 3)	PSEN1	5663	NP_015558	NM_007319	100	207784 S at
early growth response 4	EGR4	1961	NP_001956	NM_001965	ŧ	207763 at
DKFZP586A052 DKFZP586A0522 protein 2	DKFZP586A052 2	25840	NP_054752	NM_014033	ŧ	207701 S at
zinc finger protein 304	ZNF304	57343	NP_065708	NM_020657	#	207761 at
NCKassociated protein 1	NCKAPI	10787	NP 038464	NM_013436	Đ	207/38 s at
mutY homolog (E. coli)	MUTYH	4595	NP_036354	NM_012222	4D	207727_s_at
killer cell lectinlike receptor subfamily C. membe	KLRC3	3823	NP_002252	NM_002261	4D	207/23 s at
histidine triad nucleotide binding protein	HINTI	3094	NP_005331	NM_005340	40	207/21_x_at
KIAA0980 protein	K1AA0980	22981	NP_079452	NM_025176	40	207/05 s at
leukocyte immunoglobulinlike receptor, subfamily B	LILRB2	10288	NP_005865	NM_005874	ŧ	20769/_X_at
inhibin, beta C	INHBC	3626	NP_005529	NM_005538	45	207687 at
one cut domain, family member 2	ONECUT2	9480	NP 004843	NM_004852	4D	207676_at
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11 (CHL1	II (CDE)		-				
AshChi Ala Ash/Hin box ash	DEAD/H (וואַטט	1000	NI_000711			
nypometical protein MGC4293	nypometica	MOC4293					
	gastrin	MCCASOS			NM 031304		at
misc (3. cerevisiae)	DATE CALL	CAS				4D	208138 at
ASTIRE (S commission)	ASHIDA	LASII	81887	NP 112483	NM_031206	4D	208117_s_at
ne 9 onen reading from 74	chromoson	C9orf74	81605	NP_112176	NM_030914	4D	
hypothetical protein MGC 10471	hypothetica	MGC10471	81576	NP_110445	NM_030818		
nudE nuclear distribution gene E homolog like [nudE nucle	NDELI	81060	NF_110455	000000		
family with sequence similarity 49, member A	family with	FAM49A	81553		NM 030808	45 6	
tudor domain containing 3	tudor doma	TDRD3	81550		NM 030794		208097 s at
dystrophin (muscular dystrophy, Duchenne and Becke	dystrophin Becke	DMD	1756	NF_004005	MM 000014		
eta 6	integrin, beta 6	ITGB6	3694	NP_000879	NM 00488		l
eta 6	integrin, beta 6	ITGB6	3694	NP_000879	NM_000888		a
					NM_0138//		
guanylate cyclase 2F, retinal	guanylate	GUCY2F	2986	NP_001513	NM_001522		
NGFIA binding protein 1 (EGR1 binding protein 1)	NGFIA bi	NAB1	4664	NP_005957	NM_005966		208047_s_at
The second secon					NM_013303		208042 at
regulatory factor X. 2 (influences HI A class II as	regulatory	RFX2	5990	NP_000626	NM_000635		206031 S at
glutathione peroxidase 5 (epididymal androgenrelat	glutathione pe androgenrelat	GPX5	2880	NP_003987	NM_003996		200021
Ctype lectin domain family 4, member M	Ctype lect	CLEC4M	10332	NP_055072	NM_014257		208028 8 24
acrosomal vesicle protein [acrosoma	ACRVI	56	NP_064498	NM_020113		207005 X at
acrosomal vesicle protein	acrosoma	ACRV1	56	NP_064499	NM_020114		207901 x at
CD8 antigen, beta polypeptide 1 (p37)	CD8 antig	CD8B1	926	NP_004922	NM_004931		207979 S at
acrosomal vesicle protein [acrosoma	ACRV1	56	NP_064495	NM 020110		207973 X at
acrosomal vesicle protein 1	acrosoma	ACRV1	56	NP_064494	NM 020109		207072 x at
golgi apparatus protein 1	golgi appa	GLG1	2734	NP 036333	NM_012201		207960 S at
casein kinase 1, delta	casein kin	CSNK1D	1453	NP 001884	NM_001893		207945 s at
6phosphofructo2kinase/fructose2,6biphosphatase	6phospho 2	PFKFB2	5208	NP_006203	NM_006212		ZOTOLE S_II
x gene 8	paired box gene 8	PAX8	7849	NP_054698	NM_013992		207921 X BI

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Clusterin (complement lusis inhihitate Chara to	CLU	1191	NP 976084	NM 203339	ť	200/91 31
polymerase I and transcript release factor	PTRF	284119	NP 036364	7577 DAIN	5 6	208701 01
mitochondrial ribosomal protein L3	دن	11222	NP 035354	SOZZOO MIN	5 6	208789 at
microtubuleassociated protein 1 light chain 3 beta	СЗВ	01031	001000 div	NM 007208	45	208787 at
illiciolubuleassociated protein 1 light chain 3 beta		01631	ND 073770	NM 022818	4D	208786 s at
uopilootasuympno	MADII Cab	81631	NP 073729	NM 022818	4D	208785 s at
membrane cofactor protein (CD46,	MCP	4179	NP_/58865	NM_1/2333	ŧ	20070J_3 at
follistatinlike 1	FSTL1	11167	NP 009016	COOLDO MINI	3 6	208783 6 04
discoidin domain receptor family, member 1		/80	NF 034099	MINI OCCOS	j (208782 at
ribosomal protein L22	2	6146	NP 0009/4	NIM 012002	4 6	208779 x at
heterogeneous nuclear ribonucleoprotein R	^	10236	NF 00007	NIM 000000	40	208768 v at
delta sleep inducing peptide, immunoreactor		1831	NP 004080	NM 005826	4) 6	208766 s at
ATP synthase, H+ transporting, mitochondrial F0 co	ATP5L	10632	NP_006467	NM_0004/6	ŧ	200740_x_at
sin3associated polypeptide, 18kDa	SAP18	10284	NP_005861	0/8C00 IMN	ŧ	200742 3 01
sin3associated polypeptide, 18kDa	SAP18	10284	NP_005861	NM 005870	ŧ	208747 s at
RAB2, member RAS oncogene family	RAB2	5862	0C8200 AN	C00200 TAINI	3 6	208740 at
cyclin D1 (PRAD1: parathyroid adenomatosis 1)	CCNDI	595	NP 444284	NM 003005	3 8	208734 × at
ribosomal protein L39	RPL39	6170	NP_000991	000000 MIN	3 6	208711 s at
peroxiredoxin l	PRDX1	5052	NP_002565	1 C200 IAIN	đ	208605 5 24
proliferationassociated 2G4, 38kDa	PA2G4	5036	NP_000182	161000 TAIN	ð	208680 at
dolichyldiphosphooligosaccharideprotein glycosyltr	DDOST	1650	NP_005207	NW 00210	3 6	208676 s at
ribosomal protein \$14	RPS14	6208	NP 005608	/19C00 IMM	à	208674 v at
thioredoxin domain containing 7 (protein disulfide	TXNDC7	10130	NP 005/33	MM 005/42	4 6	208646 at
leukocyte immunoglobulinlike receptor, subfamily	LILRB6	79168	NP_077294	NM_024318	à é	208638 at
corticotropin releasing hormone receptor 1	CRHRI	1394	NP 004373	NM 004382	ŧ	10 V CCC007
H2B histone family, member S	H2BFS	54145	NP_059141	NM 01/445	ð	208802 v at
histone 1, H3c	HIST1H3C	8352	NP_003522	NM 003531	ŧ	200577 at
acidic (leucinerich) nuclear phosphoprotein 32 fam	ANP32D	23319	NF_050556	012404		
interferon, alpha 6	IFNA6	3443	NP 066282	700120 MIN	40 8	208571 at
histone I, H2bh	HIST1H2BH	8345	NP 003515	NM 031003	40	208548 at
histone I, H2be	nio i inzBE	thr.Co	11000011		5	20000

keratin 8	KRT8	3856	NP_002264	NM 002273	4D	209008_x_at
NPD014 protein	NPD014	57035	NP 996918	NM_207035	4D	209006 s at
Fbox and leucinerich repeat protein 5	FBXL5	26234	NP_036293	NM_012161	4D	209004 s at
anaphase promoting complex subunit 13	ANAPC13	25847	NP_056206	NM_015391	4D	209001 s_at
transcription factor 12 (EITF4, helixloophelix tran	TCF12	6938	NP_996921	NM_207038	4D	208986_at
uroporphyrinogen decarboxylase	UROD	7389	NP_000365	NM_000374	40	2089/0 s at
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex	NDUFA9	4704	NP_004993	NM_005002	ŧ	200909_at
Kruppellike factor 6	KLF6	1316	NP 001291	NM_001300	45	208961 s at
Kruppellike factor 6	KLF6	1316	NP_001291	NM_001300	4	208960 s at
thioredoxin domain containing 4 (endoplasmic retic	TXNDC4	23071	NP_055866	NM_015051	40	208957_at
protein phosphatase 4 (formerly X), catalytic subu	PPP4C	5531	NP 002711	NM_002720	4D	208932_at
ring finger protein [1]	RNFII	26994	NP_055187	NM_014372	4D	208924 at
NAD kinase	FLJ13052	65220	NP_075394	NM_023018	4D	208917 x at
golgi associated, gamma adaptin ear containing, AR	GGA2	23062	NP_619581	NM_138640	45	208914_at
golgi associated, gamma adaptin ear containing, AR	GGA2	23062	NP_055859	NM_015044	ŧ	208915_at
ribosomal protein S28	RPS28	6234	NP_001022	NM_001031	45	208903 at
DEAD (AspGluAlaAsp) box polypeptide 18	DDX18	8886	NP_006764	NM_006773	40	208896_at
DEAD (AspGluAlaAsp) box polypeptide 18	DDX18	8886	NP_006764	NM_006773	4D	208895 s at
	DUSP6	1848	NP_001937	NM_001946	4D	208892 s at
chromosome 5 open reading frame 18	C5orf18	7905	NP_005660	NM_005669	40	208873 s at
thioredoxin	NXT	7295	NP_003320	NM_003329	4D	208864 s at
splicing factor, arginine/serinerich (splicing f	SFRS1	6426	NP_008855	NM_006924	4D	208863_s_at
cisplatin resistanceassociated overexpressed prote-	CROP	51747	NP_006098	NM_006107	4D	208835 s at
ribosomal protein L23a	RPL23A	6147	NP_000975	NM_000984	4D	208834 x at
ataxin 10	ATXN10	25814	NP_037368	NM_013236	40	208855 s at
polymerase (DNA directed), epsilon 3 (p17 subunit)	POLE3	54107	NP_059139	NM_017443	45	208828_at
proteasome (prosome, macropain) subunit, beta type	PSMB6	5694	NP_002789	NM_002/98	ŧ	7000Z/_at
histidine triad nucleotide binding protein 1	HINTI	3094	NP_005331	NM_005340	45	208826 x at
ribosomal protein L23a	RPL23A	6147	NP 000975	NM_000984	4D	208825_x_at

C2f protein	C2F	10436	NP_006322	NM_006331	4D	209233_at
WD repeat domain 45	WDR45	11152	NP_009006	NM_007075	4D	209217_s_at
Kruppellike factor 5 (intestinal)	KLF5	688	NP_001721	NM 001730	40	209211 at
downregulator of transcription 1, TBPbinding (nega	DR1	1810	NP_001929	NM_001938	ŧ	209188_X_at
downregulator of transcription 1, TBPbinding (nega	DRI	1810	NP_001929	NM_001938	ŧ	200180 at
insulin receptor substrate 2	IRS2	8660		NM_003749	4	209185 S at
insulin receptor substrate 2	IRS2	8660		NM_003749	45	209184 s at
leukocyte receptor cluster (LRC) member 4	LENG4	79143	NP_077274	NM 024298	45	2091/9 s at
apoptosis antagonizing transcription factor	AATF	26574	NP_036270	NM_012138	4D	209165 at
pleckstrin homology, Sec7 and coiledcoil domains 2	PSCD2	9266	NP_059431	NM_017457	4	209138_s_at
transcription factor 3 (E2A immunoglobulin enhance	TCF3	6929		NM_003200	# #	209152_s_at
transmembrane 9 superfamily member 1	TM9SF1	10548	NP_006396	NM_006405	4D	209149 s at
corebinding factor, runt domain, alpha subunit 2;	CBFA2T2	9139	NP_005084	NM 005093	4D	209144 s at
synaptosomalassociated protein, 23kDa	SNAP23	8773	NP_003816	NM_003825	45	209130_at
squamous cell carcinoma antigen recognised by T	SART3	9733		NM_014706	ŧ	209126_S_at
tubulin, alpha 3	TUBA3	7846		NM_006009	ŧ	209118 s at
tetraspan 1	TSPANI	10103	NP_005718	NM_005727	40	209114_at
nuclear receptor coactivator 1	NCOAI	8648	NP_671756	NM_147223	6	209106_at
interferonrelated developmental regulator 2	IFRD2	7866	NP_006755	NM_006764	40	209100_at
jagged I (Alagille syndrome)	JAGI	182	NP_000205	NM_000214	45	209099_x_at
RAB5A, member RAS oncogene family	RAB5A	5868	NP_004153	NM_004162	4D	209089_at
collagen, type XVIII, alpha 1	COL18A1	80781	NP_085059	NM_030582	4D	209081_s_at
H3 histone, family 3B (H3.3B)	нзғзв	3021	NP_005315	NM_005324	4D	209069 s at
ubiquinoleytochrome c reductase binding protein	UQCRB	7381	NP_006285	NM 006294	4D	209065_at
protein kinase C binding protein 1	PRKCBP1	23613	NP 898869	NM_183048	4D	209048 s at
Synaptotagmin binding, cytoplasmic RNA interacting	SYNCRIP	10492	NP_006363	NM_006372	40	209025_s_at
KIAA0652 gene product	KIAA0652	9776	NP_055556	NM_014741	4D	209021 x at
PTEN induced putative kinase 1	PINK1	65018	NP_115785	NM_032409	4D	209019 s at
DnaJ (Hsp40) homolog, subfamily B, member 6	DNAJB6	10049	NP 490647	NM_058246	4D	209015_s_at

DNA segment on chromosome 4 (unique) 23.1	D4S234E	27065	NP 055207	NM 014392	40	70506 X at
RRS1 ribosome biogenesis regulator homolog (S. cer	RRS1	23212		NM_OLDINA	ŧ	
hypothetical protein MGC11061	MGC11061	84272		NM 032312	ŧ	
exosome component 2	EXOSC2	23404	NP_055100	INM 014285	5 5	
ash2 (absent, small, or homeotic)like (Drosophila)	ASH2L		NP 004665	NM 0046/4) é	a
hydroxysteroid dehydrogenase like 2	HSDL2	5	NP_115679	NM 032303	ŧ	
hydroxysteroid dehydrogenase like 2	HSDL2	84263	NP_115679	NM_032303	5	
polymerase (RNA) II (DNA directed) polypeptide	POLR2F	5435	NP_068809	NM_0219/4	÷	E
RNA binding motif protein 30	RBM30	83759	NP_113680	NM 031492	ŧ	a
chromosome open reading frame 48	C1orf48	25936	NP_056286	NM_015471	4D	209484 s_at
ubiquitin specific protease 15	USP15	9958	NP_006304	NM_006313	4D	2094/5 at
TRAF family memberassociated NFKB activator	TANK		NP_004171	NM_004180	4D	209451_at
Osialoglycoprotein endopentidase	OSGEP	55644	NP_060277	NM_017807	4D	209450_at
phosphoribosyl pyrophosphate amidateous face	PPAT	5471	NP 002694	NM_002703	40	209434 s at
alphamethylacylCoA racemase	AMACR	23600	NP_055139	NM_014324	45	209426 S at
chromosome 22 open reading frame 10	C22orf19	8563	NP_001002878	NM_001002878	£	209418 S at
transmembrane protein	TMEMI	7109	NP_003265	NM_003274	ŧ	209412 81
growth factor receptorbound protein 10	GRB10	2887	NP_001001550	NM_001001550	ŧ	200412 S at
growth factor receptorbound protein 10	GRB10	2887	NP_001001549	NM_001001549	ŧ	200410 at
cytochrome b5	CYB5	1528	NP_001905	NM_001914	45	209366 x at
RAD50 homolog (S. cerevisiae)	RAD50	10111	NP_597816	NM_133482	ŧ	209349 at
MYC associated factor X	MAX	4149	NP 660088	NM_145113	ŧ	209331 S at
hypothetical protein MGC2198	MGC2198	192286	NP_620175	NM_138820	4	209328 x at
polymerase (RNA) I polymentide C 30kDa	POLRIC	9533	NP_976035	NM_203290	ŧ	20931/ at
BCL2/adenovirus E1B 19kDa interacting provide	BNIP2	663	NP_004321	NM_004330	40	209508 s at
growth arrest and DNAdamageinducible beta	GADD45B	4616	NP_056490	NM 015675	ŧ	200208 - 1
carbonic anhydrase II	CA2	760	NP_000058	NM_00006/	ŧ	200206
glutaredoxin (thioltransferase)	GLRX	2745	NP_002055	NM_002064	ŧ	200201 at
vacuolar protein sorting 45A (veast)	VPS45A	11311	NP 009190	NM_007259	ŧ	200276 at
methyltransferase like 3	METTL3	56339	NP_062826	NM_019852	ŧ	200268 at
tubulin alpha 6	TUBA6	84790	NP_116093	NM 032704	ŧ	3000CE 30000
chloride channel 7	CLCN7	1186	NP_001278	NM_001287	ŧ	200251 at

ATPase Ca++ transporting type 2C mamber 1	ATP2C1	27032	NP 001001487	NM 001001487	4D	209934 s at
CD300A antigen	CD300A	11314	NP_009192	NM_007261	40	209933 s at
histone I, H2bd	HIST1H2BD	3017	NP_619790	NM_138720	4D	209911 x_at
ataxia telangiectasia and Rad3 related	ATR	545	NP_001175	NM_001184	Ð	209902 at
SEC31like 2 (S. cerevisiae)	SEC31L2	25956	NP_937781	NM_198138	4D	209889 at
Raslike without CAAX 1	RITI	6016	NP_008843	NM_006912	4D	209882 at
linker for activation of T cells	LAT	27040	NP_055202	NM_014387	4D	209881 s at
selectin P ligand	SELPLG	6404	NP_002997	NM_003006	40	209880 s at
synuclein, gamma (breast cancerspecific protein 1)	SNCG	6623	NP_003078	NM_003087	45	2098/7_at
adrenergic, alpha2A, receptor	ADRA2A	150	NP_000672	NM_000681	4D	209869 at
latrophilin 3	LPHN3	23284	NP 056051	NM_015236	4D	209867 s at
frequently rearranged in advanced Tcell lymphomas	FRAT2	23401	NP_036215	NM_012085	ŧ	207004_at
tumor protein p73like	TP73L	8626	NP_003713	NM_003722	ŧ	209863 S at
proteasome (prosome, macropain) activator subunit	PSME3	10197	NF_/89839	NV 170803	ŧ	an e coord
CDC42 effector protein (Rho GTPase binding) 2	CDC42EP2	10435	NP_006770	NM 176863	ŧ	209030 S at
leucine rich repeat neuronal 3	LRRN3	54674	NP_060804	NM_018334	ŧ	209841 s at
leucine rich repeat neuronal 3	LRRN3	54674	NP_060804	NM_018334	1	209840 s at
palmitoylprotein thioesterase 2	PPT2	9374	NP_620312	NM 138934	ŧ	209826 at
jagged 2	JAG2	3714	NP_002217	NM_002226	ŧ	209784 s at
CD24 antigen (small cell lung carcinoma cluster 4	CD24	934	NP_037362	NM_013230	4	209772 s at
itchy homolog E3 ubiquitin protein ligase (mouse)	ITCH	83737	NP_113671	NM_031483	40	209744_x_at
itchy homolog E3 ubiquitin protein ligase (mouse)	ITCH	83737	NP_113671	NM_031483	4D	209743 s at
Ctype lectin domain family 2, member B	CLEC2B	9976	NP_005118	NM_005127	£	209732 at
solute carrier family 35 (UDPglucuronic acid/UDPNa	SLC35D1	23169	NP_055954	NM_015139	ŧ	209/11_at
fatso	FTO	79068	XP_051200	XM 051200	ŧ	209702 at
S100 calcium binding protein, beta (neural)	S100B	6285	NP_006263	NM_006272	4	209686_at
carboxylesterase 2 (intestine, liver)	CES2	8824	NP_932327	NM_198061	40	20966/_at
CDC16 cell division cycle 16 homolog (S. cerevisia	CDC16	888	NP_003894	NM_003903	ŧ	207039 S at
promyelocytic leukemia	PML	5371	NP 150241	NM 053238	ŧ	209050 at
regulator of Gprotein signalling 12	RGS12	6002	NP 937870	NM 198227	ŧ	209039 S at
cybresed						20000

FK506 binding protein 1A, 12kDa	FKBP1A	2280	NP 000792	NM_000801	4D	210186_s_at
tolllike receptor 1	TLR1	7096	NP_003254	NM_003263	4D	210176 at
protein tyrosine phosphatase, receptor type, J	PTPRJ	5795	NP_002834	NM_002843	4D	210173_at
plateletactivating factor acetylhydrolase, isoform	PAFAH1B2	5049	NP_002563	NM_002572	4D	210160_at
leukocyte immunoglobulinlike receptor, subfamily B	LILRB4	11006	NP_006838	NM_006847	4D	210152_at
dCMP deaminase	DCTD	1635	NP_001912	NM_001921	4D	210137_s_at
leukotriene B4 receptor	LTB4R	1241	NP_858043	NM_181657	4D	210128_s_at
barrier to autointegration factor 1	BANFI	8815	NP_003851	NM_003860	4D	210125_s_at
potassium inwardlyrectifying channel, subfamily J,	KCNJ15	3772	NP_002234	NM_002243	4D	210119_at
sperm associated antigen 1	SPAG1	6674		NM_003114	4D	210117_at
inversin	INVS	27130	NP_899068	NM_183245	4D	210114_at
nucleolar protein 7, 27kDa	NOL7	51406	NP_057251	NM_016167	4D	210097_s_at
par3 partitioning defective 3 homolog (C. elegans)	PARD3	56288	NP_062565	NM_019619	4D	210094 s at
magonashi homolog, proliferationassociated (Drosop	MAGOH	4116	NP_002361	NM_002370	đĐ	210093_s_at
magonashi homolog, proliferationassociated (Drosop	MAGOH	4116		NM_002370	4D	210092_at
myosin, light polypeptide 4, alkali; atrial, embry	MYL4	4635	NP_001002841	NM_001002841	4D	210088_x_at
ribosomal protein L5	RPL5	6125	NP 000960	NM_000969	4D	210034_s_at
APEX nuclease (multifunctional DNA repair enzyme)	APEX1	328	NP_542380	NM_080649	4D	210027_s_at
polycomb group ring finger 1	PCGF1	84759		NM_032673	4D	210023 s at
mucosa associated lymphoid tissue lymphoma translo	MALTI	10892	NP_006776	NM_006785	4D	210017_at
hemopexin	HPX	3263	NP_000604	NM_000613	4D	210013_at
solute carrier family 25 (mitochondrial carrier; c	SLC25A1	6576	NP_005975	NM_005984	4D	210010_s_at
oxidised low density lipoprotein (lectinlike) rece	OLRI	4973	NP_002534	NM_002543	4D	210004_at
suppressor of cytokine signaling 1	SOCS1	8651	NP_003736	NM_003745	4D	209999 x at
zinc finger protein 268	ZNF268	10795	NP_003406	NM_003415	4D	209989_at
jumonji domain containing 2C	JMJD2C	23081	NP_055876	NM_015061	4D	209984_at
lipoprotein, Lp(a)	LPA	4018	NP_005568	NM_005577	4D	209978 s_at
JTV1 gene	JTVI	7965	NP_006294	NM_006303	4D	209971 x at
melanoma antigen family A, 3	MAGEA3	4102	NP_005353	NM_005362	4D	209942 x at

	DOLLAR!	5710	NP 722544	NM 153822	4D	210459 at
chromosome 10 open reading frame 28	C10orf28	27291	NP_055287	NM_014472	4D	210455_at
co						
ATB arthur III	ATDSI NI4	10632	NP 006467	NM 006476	45	210453 x at
ribonucleoprot	MADVIA	1/33	ND 001306	NM 001315	40	210449 x at
Sjogren syndrome antigen A2 (60kDa	SSA2	6738	NP_004591	NM_004600	4D	210438 x at
defensin, beta 1	DEFB1	1672	NP_005209	NM_005218	4D	210397 at
myosin, light polypeptide 4, alkali; atrial, embry	MYL4	4635	NP_002467	NM_002476	4D	210395_x_at
synovial sarcoma, X breakpoint 4	SSX4	6759	NP_005627	NM_005636	4D	210394_x_at
histone 1, H2bg	HIST1H2BG	8339	NP_003509	NM_003518	4D	210387_at
calcium channel, voltagedependent, alpha 1G subuni	CACNAIG	8913	NP_938198	NM_198384	ŧ	210580_S_at
lymphocyte antigen 9	LY9	4063	NP_002339	NM_002348	ŧ	210370 s at
Bcell CLL/lymphoma 11A (zinc finger protein)	BCLIIA	53335	NP_060484	NM_018014	4D	210347 s at
myelin transcription factor I	MYTI	4661	NP_004526	NM_004535	4D	210341_at
peptidylglycine alphaamidating monooxygenase COOHt	PAMCI	9182	NP_005438	NM_005447	45	210335_at
BTB/POZ KELCH domain protein	ENC2	64410	NP_071925	NM_022480	4D	210307 s at
l(3)mbtlike (Drosophila)	L3MBTL	26013	NP_056293	NM_015478	4D	210306_at
mab21like 2 (C. elegans)	MAB21L2	10586	NP_006430	NM_006439	4D	210303 at
xanthine dehydrogenase	XDH	7498	NP_000370	NM_000379	4D	210301 at
mitogenactivated protein kinase kinase kinase 7 in	MAP3K7IP2	23118	NP_055908	NM_015093	4D	210284 s at
G proteincoupled receptor 18	GPR18	2841	NP_005283	NM_005292	4D	210279_at
cytochrome P450, family 2, subfamily B, polypeptid	CYP2B6	1555	NP_000758	NM_000767	40	210272_at
neurogenic differentiation 2	NEUROD2	4761	NP_006151	NM_006160	4D	210271_at
membranespanning 4domains, subfamily A, member 3 (MS4A3	932	NP_006129	NM_006138	ŧ	210254_at
adenylosuccinate lyase	ADSL	158	NP_000017	NM_000026	10	210250 x at
cyclindependent kinase inhibitor 2D (p19, inhibits	CDKN2D	1032	NP_524145	NM 079421	4D	210240_s_at
protein tyrosine phosphatase, receptor type, f pol	PPFIAI	8500	NP_803172	NM_177423	4D	210236_at
interleukin 1 receptor accessory protein	ILIRAP	3556	NP 608273	NM_134470	4D	210233_at
transferrin receptor 2	TFR2	7036	NP_003218	NM_003227	4D	210215_at
syntaxin 11	SIXII	86/6	NP_005/55	107 COO TAIN	+	710190 at

						nonAT
210495_x_at	4D	NM_212474	NP_997639	2335	FNI	fibronectin 1
210506_at	4D		NP 004470	2529	FUT7	fucosyltransferase 7 (alpha (1,3) fucosyltransfera
210546_x_at	4D		NP_001318	1485	CTAGIB	cancer/testis antigen IB
210550 s_at	4D		NP 002882	5923	RASGRFI	Ras proteinspecific guanine nucleotidereleasing fa
210594 x at	4D	NM_003953	NP_003944	9019	MPZLI	myelin protein zerolike 1
210613_s_at	4D	NM_145738	NP_663791	9145	SYNGRI	synaptogyrin 1
210617_at	4D	NM_000444	NP_000435	5251	PHEX	phosphate regulating endopeptidase homolog, Xlinke
210645_s_at	4D	NM_003316	NP_003307	7267	TTC3	tetratricopeptide repeat domain 3
210646_x_at	4D	NM_012423	NP_036555	23521	RPL13A	ribosomal protein L13a
210681_s_at	4D	NM_006313	NP_006304	9958	USP15	ubiquitin specific protease 15
210695 s at	4D	NM_016373	NP_057457	51741	WWOX	WW domain containing oxidoreductase
210724_at	4D	NM_032571	NP_115960	84658	EMR3	egflike module containing, mucinlike, hormone rece
210734_x_at	4D	NM_002382	NP_002373	4149	MAX	MYC associated factor X
210740 s at				3705	IJPKI	inositol 1,3,4triphosphate 5/6 kinase
210749 x at	40		NP_001945	780	DDR1	discoidin domain receptor family, member 1
210750 s at	45	NM_001003809	NP_001003809	9229	DLGAP1	discs, large (Drosophila) homologassociated protei
210754_s_at	4D	NM_002350	NP_002341	4067	LYN	vyes I Yamaguchi sarcoma viral related oncogene hom
210772_at	4b	NM_001005738	NP_001005738	2358	FPRL1	formyl peptide receptorlike 1
210774 s at	4D		NP_005428	8031	NCOA4	nuclear receptor coactivator 4
210775_x_at	4D	NM_001229	NP_001220	842	CASP9	caspase 9, apoptosisrelated cysteine protease
210779_x_at	4D			8487	SIPI	survival of motor neuron protein interacting prote
210787_s_at	4D		NP_757380	10645	CAMKK2	calcium/calmodulindependent protein kinase kinase
210789_x_at	4D		NP_001806	1084	CEACAM3	carcinoembryonic antigenrelated cell adhesion mole
210792 x at	4D	NM_006427	NP 006418	10572	SIVA	CD27binding (Siva) protein
210811 s at				54555	DDX49	DEAD (AspGluAlaAsp) box polypeptide 49
210818_s_at	4D	NM_001186	NP_001177	571	BACHI	BTB and CNC homology 1, basic leucine zipper trans
210826_x_at			NP_579916	5884	RADI7	RAD17 homolog (S. pombe)
210886_x_at	45	NM_007233	NP_009164	11257	TP53AP1	TP53 activated protein 1

APPENDIX

ubiquitin protein ligase E3A (human papilloma	UBE3A	7337	NP_570853	NM_130838	4D	211285 s at
nuclear transcription factor Y, ganuma	NFYC	4802	NP_055038	NM_014223	4D	211251_x_at
SH3domain binding protein 2	SH3BP2	6452	NP_003014	NM 003023	4D	211250 s at
cytochrome P450, family 4, subfamily A, polypeptid	CYP4A11	1579	NP_000769	NM_000778	4D	211231_x_at
tumor necrosis factor receptor superfamily, member	TNFRSF10C	8794	NP_003832	NM_003841	4D	211163_s_at
actinin, alpha l	ACTNI	87	NP 001093	NM_001102	4D	211160 x at
NGFIA binding protein 1 (EGR1 binding protein 1)	NABI	4664	NP_005957	NM_005966	4D	211139_s_at
leukocyte immunoglobulinlike receptor, subfamily	LILRB3	11025	NP_006855	NM_006864	4D	211135_x_at
ectodysplasin A	EDA	1896	NP_001005611	NM_001005611	4D	211128 at
survival of motor neuron protein interacting prote	SIPI	8487	NP 003607	NM_003616	4D	211115 x at
survival of motor neuron protein interacting prote	SIPI	8487	NP_003607	NM_003616	Đ	211114 x at
leukocyte immunoglobulinlike receptor, subfamily	LILRA2	11027	NP_006857	NM_006866	4D	211102 s_at
preBcell leukemia transcription factor 2	PBX2	5089	NP_002577	NM_002586	4D	211097_s_at
diazepam binding inhibitor (GABA receptor modulato	DBI	1622	NP_065438	NM_020548	4D	211070_x_at
exostoses (multiple)like 3	EXTL3	2137	NP_001431	NM_001440	4D	211051_s_at
peroxisomal biogenesis factor 7	PEX7	5191	NP_000279	NM_000288	4D	211033_s_at
aldehyde dehydrogenase 3 family, member B1	ALDH3B1	221	NP_000685	NM_000694	4D	211004 s at
growth factor receptorbound protein 10	GRB10	2887	NP_001001555	NM_001001555	4D	210999 s_at
tripartite motifcontaining 23	TRIM23	373	NP_150231	NM_033228	4D	210995_s_at
Nacylsphingosine amidohydrolase (acid ceramidase)	ASAHI	427	NP_004306	NM_004315	\$	210980_s_at
FAST kinase	FASTK	10922	NP 148936	NM_033015	4D	210975 x at
KIAA0669 gene product	KIAA0669	9819	NP_055594	NM_014779	4D	210954 s at
insulin promoter factor 1, homeodomain transcripti	IPFI	3651	NP_000200	NM_000209	#	210937_s_at
ring finger protein (C3H2C3 type) 6	RNF6	6049	NP_898866	NM 183045	4D	210931_at
similar to FKSG30	LOC440915	440915	XP 496601	XM_496601	4D	210926_at
programmed cell death 10	PDCD10	11235	NP 009148	NM_007217	4D	210907_s_at
interleukin 13 receptor, alpha 1	ILI3RAI	3597	NP_001551	NM_001560	4D	210904 s at

						VIII
211298 s at	4D	NM_000477	NP_000468	213	ALB	albumin
211330 s at			NP 000401	3077	HFE	hemochromatosis
211272 x at		NM_002303	NP 002294	3953	LEPR	leptin receptor
211272 8 81		NM_004633	NP 004624	7850	IL1R2	interleukin I receptor, type []
211378 x at		NM 203431	NP 982255	5478	PPIA	peptidylprolyl isomerase A (evelophilin A)
211396_at		NM_001005412	NP 001005412	9103	FCGR2C	Fe fragment of left, low affinity He recenter for
211413 s at		NM_012387	NP_036519	23569	PADI4	peptidyl arginine deiminase type IV
211425 x at	4D	NM_005636	NP_005627	6759	SSX4	synovial sarcoma X break point 4
211433 x at	4D	NM_025182	NP_079458	80256	KIAA1539	KIAA1539
211501_s_at	4D	NM_003751	NP_003742	8662	EIF3S9	eukarvotic translation initiation factor 2
211507 s at	4D	NM_153051	NP 694691	8897	MTMR3	myofibularin related protein 2
211521_s_at	4D	NM_013385	NP_037517	27128	PSCD4	pleckstrin homology, Sec7 and coiledcoil domains
211530_x_at	4D	NM_002127	NP 002118	3135	HLAG	HI AG histogramatikilis
211531_x_at	4D		NP_955386	1	PRB1	prolinerich protein BetN1 subfamily 1
211534_x_at	4D	NM_002847	NP_002838	5799	PTPRN2	protein tyrosine phosphatase receptor type N pol
211538 s at		NM_021979	NP 068814	3306	HSPA2	heat shock 70kDa protein 2
211342 x at		NM_001014	NP_001005	6204	RPS10	ribosomal protein \$10
211548_s_at		NM_000860	NP_000851	3248	HPGD	hydroxyprostaglandin dehydrogenase 15(N/A/D)
211005 s at			NP_000848	2983	GUCY1B3	guanylate cyclase I, soluble, beta 3
211561 x at			NP_001306	1432	MAPK14	mitogenactivated protein kinase 14
211500 x_at				9577	BRE	brain and reproductive organexpressed (TNFRSF1A mo
2113/4 S at			NP_758866	4179	MCP	membrane cofactor protein (CD46, trophoblastlympho
				65005	MRPL9	mitochondrial ribosomal protein L9
				3597	IL13RA1	interleukin 13 receptor, alpha 1
=				3912	LAMB1	laminin, beta 1
	45	NM_002483	NP_002474	4680	CEACAM6	carcinoembryonic antigenrelated cell adhesion mole
		NM_000416	NP_000407	3459	IFNGR1	interferon gamma recentor i
		NM_032509	NP_115898	84549	RBM13	RNA binding motif protein 13
a		NM_014642	NP_055457	9657		IQ motif containing B1
211/11 S at	40	NM_000314	NP 000305	9073	PTEN	

myeloid/lymphoid or mixedlineage leukemia	MLL	4297	NP_005924	NM 005933	40	212076 at
transient receptor potential cation channel, subfa	TRPC4AP	26133	NP_056453	NM 015638	40	8
ring finger protein 167	RNF167	26001	NP_056343	NM_015528	ŧ	
exocyst complex component 7	EXOC7	23265	NP_056034	NM_015219	ŧ	212047 S at
antigen identified by monoclonal antibody Ki67	MK167		NP_002408	NM 002417	ŧ	212021 S at
CD44 antigen (homing function and Indian blood gro	CD44	960	390	NMI_001001390	ŧ	10011 - 10010
H3 histone, family 3B (H3.3B)	H3F3B	_		NM 001324	ŧ	212014 x at
major histocompatibility complex, class II, DP alp	HLADPAI	3113	NP_291032	NM 033554	ŧ	211000 at
collagen, type IV, alpha 1	COL4A1	1282	NP_001836	NM 001845	£	211900 at
peptidylprolyl isomerase A (cyclophilin A)	PPIA	5478	NP_066953	NM_021130	45	2119/8 X at
heat shock protein 90kDa protein 1 <u>alpha</u> (cytosolic), class A member 1	HSPCA HSP90AAI	3320	NP_005339	NM_000348	ŧ	211079 m
collagen, type IV, alpha 2	COL4A2		NP_001837	NM_001846	ŧ.	211960 of
plasminogen activator, urokinase receptor	PLAUR	5329	NP_001005377	NM_001005377	ŧ	211924 S at
prostaglandin E receptor 3 (subtype EP3)	PTGER3	5733	NP_942007	NM_198714	ŧ	211909 x at
fucosyltransferase 6 (alpha (1,3) fucosyltransfera	FUT6	2528	NP_000141	NM_000150	40	211682 X at
carcinoembryonic antigenrelated cell adhesion mole	CEACAMI	634	NP_001703	NM 001/12	ŧ	21 1000 A at
carcinoembryonic antigenrelated cell adhesion mole	CEACAM7	1087	NF_000021	NINI_000850	9 6	211002
trithor	TILL L		ND COORD		45	211848 s at
cyclin EZ	CCME?		NP 005926	NM 005935	4D	211826 s at
5,	COMES	0134	NP 477097	NM 057749	4D	211814 s at
potassium inwardlyrectifying channel, subfamily	KCNJ15	3772	NP_733933	NM_170737	40	211806 s at
peptidylprolyl isomerase A (cyclophilin A)	PPIA	5478	NP_066953	NM 021130	ŧ	21100 x at
ubiquitinconjugating enzyme E2B (RAD6 homolog)	UBE2B	7320	NP_003328	NM_003337	ŧ	211765
tubulin alpha 6	TUBA6	84790	NP_116093	NM 052/04	ŧ	211762 au
elastase 3A, pancreatic (protease E)	ELA3A	10136	NP_005738	NM 005747	45	211750 x at
Fc fragment of IgE, high affinity I, receptor for:	FCERIA	2205	NP_001992	NM 002001	ŧ	211734 S at
synovial sarcoma, X breakpoint 3	SSX3	10214	NP_783642	NM_175711	ŧ	211/31 x at
BH3 interacting domain death agonist	BID	637	NP 932071	NM 197967	ŧ	211721 s at
multipl						

U3 snoRNP protein 4 homolog	IMP4	92856	NP_219484	NM 033416	40	212411_at
KIAA0082	K1AA0082	23070	NP 055865			
phosphoribosylglycinamide formyltransferase, phosp	GART					
ATPase, Ca++ transporting, cardiac muscle, slow tw	ATP2A2	488	NP_001672) g
KIAA0323	KIAA0323	23351	NP 056114			
amine oxidase (flavin containing) domain 2	AOF2	23028				al
myosin ID	MYOID	4642	NP_056009	NM_015194	40	
glucosamine (Nacetyl)6sulfatase (Sanfilippo diseas	GNS	2799				
DKFZP564F0522 DKFZP564F0522 protein	DKFZP564F0522	25940	NP_056290	NM_015475		
transcription factor Dp1	TFDP1	7027	NP 009042	NM_007111		
KIAA0746 protein	KIAA0746	23231	NP_056002	NM_015187	4D	
proteasome (prosome, macropain) 26S subunit, nonAT	PSMD14	10213	NP_005796			
lipin I	LPINI	23175	NP_663731	NM_145693	4D	212274_at
ribosomal protein L17	RPL17	6139	NP_000976	NM_000985		at
serine (or cysteine) proteinase inhibitor, clade B	SERPINB1	1992	NP_109591	NM_030666		
dystonin	DST	667	NP_899236	NM_183380	40	212254 s at
tubulin, alpha 1 (testis specific)	TUBAI	7277	NP_005991			212242 at
Fbox protein 21	FBXO21	23014	NP_296373	NM_033624		212229 s at
subunit						
professine (procome macronain) activate	4	23198	NP 055429	NM 014614	4D	212222_at
optic atrophy 1 (autosomal dominant)	OPA1	4976	NP_570845	NM_130832	4D	212213_x_at
DKFZP564G202 DKFZP564G2022 protein	DKFZP564G202	25963	NP_036312	NM_013497	45	21220+_at
interferon induced transmembrane protein 3 (18U)	IFITM3	10410	NP 066362	NM 021034		212201 at
chromosome 10 open reading frame 61	C10orf61	26123	NP_056446	NM_015631		21222 at
expressed in Teells and eosinophils in atopic derm	ETEA	23197	NP 055428	NM 014613		212123 at
expressed in Teells and eosinophils in atopic derm	ETEA	23197	NP_055428	NM 014613		212106 at
karyopherin alpha 6 (importin alpha 7)	KPNA6	23633	NP 036448	NM_012316	40	212101 at
Era Gproteinlike I (E. coli)	ERAL1	26284	NP_005693	NM 005702		21208/ s at
caldesmon l	CALDI	800	NP 149130	NM_033139		212077 at
(trithor						212022

	BICD2	23299	NP_001003800	NM_001003800	40	212/02 s at
septin 10	SEPT10	151011	NP_653311	NM 144710	ŧ	212096 S at
hypothetical protein LOC162427	LOC162427	162427	NP_835227	NM_178126	£	212699 at
protein phosphatase 1H (PP2C domain containing)	PPMIH	57460		XM_350880	4D	212686 at
zinc finger protein 3 (A851)	ZNF3	7551	NP_116313	NM 032924	40	212684 at
erythrocyte membrane protein band 4.11ike 3	EPB41L3	23136	NP_036439	NM_012307	4D	212681 at
KIAA0582	KIAA0582	23177	NP 055962	NM 015147	4	212677 s at
secreted protein, acidic, cysteinerich (osteonecti	SPARC	6678	NP_003109	NM_003118	45	212667_at
TCDDinducible poly(ADPribose) polymerase	TIPARP	25976	NP_056323	NM_015508	ŧ	212665 at
peptidylprolyl isomerase A (cyclophilin A)	PPIA	5478	NP_066953	NM_021130	45	212601 X at
interleukin 1 receptor antagonist	ILIRN	3557	NP_776215	NM_173843	40	212659 s at
sorting nexin 4	SNX4	8723	NP_003785	NM_003794	ŧ	212652 s_at
DEAH (AspGluAlaHis) box polypeptide 29	DHX29	54505	NP_061903	NM_019030	45	212648 at
protein tyrosine phosphataselike (proline instead	PTPLB	201562	NP_940684	NM_198402	4D	212640_at
quaking homolog, KH domain RNA binding (mouse)	QKI	9444	NP_996737	NM_206855	ŧ	-12000_at
mitochondrial ribosomal protein S31	MRPS31	10240	NP_005821	NM_005830	16	12627 at
protein tyrosine phosphatase, receptor type, C	PTPRC	5788	NP_563579	NM 080922	ŧ	212602 at
zinc finger protein 451	ZNF451	26036	NP_056370	NM_015555	6	212588 -t
signal transducer and activator of transcription 5	STAT5B	6777	NP_036580	NM_012448	40	212550 at
FADsynthetase	PP591	80308	NP_079483	NM_025207	45	212341_at
chromodomain helicase DNA binding protein llike	CHDIL	9557	NP_004275	NM_004284	ŧ	212339_at
ribosomal protein L17	RPL17	6139	NP_000976	NM_000985	ŧ	21253/ x at
DNA segment, Chr 15, Wayne State University 75, ex	D15Wsu75e	27351	XP_039495	AM_039495	ŧ	-12/2/at
phosphodiesterase 8A	PDE8A	5151	NP_775658	NM 173456	ŧ	212527 at
chromosome 10 open reading frame 22	C10orf22	84890	NP_116193	NM 032804	45	212500_at
KIAA0553 protein	KIAA0553	23131	NP_001002909	NM_001002909	40	212487 at
DnaJ (Hsp40) homolog, subfamily C, member 13	DNAJC13	23317	NP_776184	NM_173823	4D	212467 at
fibronectin 1	FNI	2335	NP_002017	NM_002026	4Đ	212464 s at
transcription factor binding to IGHM enhancer 3	TFE3	7030	NP_006512	NM_006521	4	212457 at
general transcription factor HIC polymentide 2	GTF3C2	2976	NP_001512	NM_001521	4D	212429 s at
septin 6	SEPT6	23157	NP_665798	NM 145799	ŧ	212417 dt

	HIC2 hypermethylated in concer	23119	NP 055909	NM 015094	40	212904 at
peptidylglycine alphaamidating monooxygenase	PAM	5066	NP_620177	NM_138822	4D	212958 x at
polymerase (RNA) II (DNA directed) polypeptide I,	POLR2I	5438		NM_006233	ŧ	212905 s_at
collagen, type VI, alpha I	COL6A1	1291	NP_001839	NM 001848	ŧ	212940 at
hypothetical protein DKFZp564D172	DKFZP564D172	83989	NP_114431	NM_032042	40	212936_at
ribosomal protein L13	RPL13	6137	NP 000968	NM_000977	4D	212933_x_at
TSPYlike 4	TSPYL4	23270	NP_067680	NM_021648	4D	212928_at
SMC5 structural maintenance of chromosomes 5like 1	SMC5L1	23137	NP_055925	NM_015110	40	212926_at
chromosome 19 open reading frame 21	C19orf21	126353	NP_775752	NM_173481	40	212925_at
SET and MYND domain containing 2	SMYD2	56950	NP_064582	NM_020197	40	212922 s_at
cleavage stimulation factor, 3' preRNA, subunit 2.	CSTF2T	23283	NP_056050	NM_015235	4D	212905_at
Mphase phosphoprotein 10 (U3 small nucleolar ribon	MPHOSPH10	10199	NP_005782	NM_005791	ŧ	212885_at
chromosome 21 open reading frame 25	C21orf25	25966	NP_950251	NM_199050	45	212675 s at
ubiquitin specific protease 49	USP49	9477	NP_004266	NM_004275	40	212872 s at
CDPdiacylglycerol synthase (phosphatidate cytidyly	CDS2	8760	NP_003809	NM 003818	ŧ	212002_at
KIAA0276 protein	KIAA0276	23142	NP_055930	NM_015115	ŧ	212853 at
Sjogren syndrome antigen A2 (60kDa, ribonucleoprot	SSA2	6738	NP_004591	NM_004600	ŧ	212659 S at
polymerase (DNAdirected), delta 3, accessory subun	POLD3	10714	NP_006582	IAC900 WN	ŧ	212020 at
EGFlikedomain, multiple 5	EGFL5	1955	XP_376905	XM_3/6905	ŧ	212626 at
PAX transcription activation domain interacting pr	PAXIPIL	22976	NP_031375	NM 007349	10	212825 at
DnaJ (Hsp40) homolog, subfamily B, member 5	DNAJBS	25822	NP 036398	NM_012266	8	212817_at
KIAA0877 protein	KIAA0877	23333	XP_371891	XM_371891	4D	212792_at
ribosomal protein L13a	RPL13A	23521	NP_036555	NM_012423	4D	212790 x_at
HDCMA18P protein	HDCMA18P	51574	NP 057732	NM_016648	4D	212785 s at
GTP_binding protein	GTP	92170	NP_612393	NM_138384	4D	212767_at
ankyrin repeat and sterile alpha motif domain cont	ANKSI	23294	NP_056060	NM_015245	4D	212747_at
pleckstrin homology domain containing, family E (w	PLEKHEI	23239	NP_919431	NM_194449	4 0	212719_at
cMpl binding protein	LOC113251	113251	NP_443111	NM 0328/9	ť	212/14 at

chromosome 14 open reading frame 124	C14orf124	56948	NP_064580	NM 020195	40	213398 S at
DEAD/H (AspGluAlaAsp/His) box polypeptide 11 (CHL1	DDXII	1663	NP_004390	NM_004399	ŧ	2133/6_8_at
caspase 8, apoptosisrelated cysteine protease	CASP8		NP_001219	NM_001228	ð	213373 s at
transmembrane and coiledcoil domains I	TMCC1	23023	NP_055823	NM_015008	4D	213349_at
KIAA0495	KIAA0495	57212	NP 997189	NM_207306	40	213340 s at
vesicleassociated membrane protein l (synaptobrevi	VAMPI	6843	NP_055046	NM_014231	45	215526_at
chromosome 6 open reading frame 162	C6orf162	57150	NP_065158	NM_020425	40	213312 at
phosphoribosylformylglycinamidine synthase (FGAR a	PFAS	5198	NP_036525	NM_012393	45	213302_at
zinc finger protein 248	ZNF248	57209	NP_066383	NM_021045	45	213269_at
zinc finger protein 187	ZNF187	7741	NP_689949	NM_152736	40	213218_at
small nuclear RNA activating complex, polypeptide	SNAPCS	10302	NP_006040	NM_006049	4	213203_at
component of oligomeric golgi complex 7	COG7	91949	NP_705831	NM_153603	4D	213190_at
Sp3 transcription factor	SP3	6670	NP_003102	NM_003111	4D	213168_at
glycine cleavage system protein H (aminomethyl car	GCSH	2653	NF_0044/4	NW 004483	ŧ	
glycine cleavage system protein H (aminomethyl	GCSH	2633	NF_0044/4	NM 004483	ē	212122 2 4
TSPYlike 5	TSPYL5	85453	NP_277047	NM 033512	ŧ	213120 s at
DnaJ (Hsp40) homolog, subfamily C, member 9	DNAJC9	23234	NP 056005	NM 015190	ŧ	213132 X at
ribosomal protein L23a	RPL23A	6147	NP 000975	NM_000984	4D	213084 x at
PLSC domain containing protein	LOC254531	254531	NP_705841	NM_153613	4D	213078 x_at
zinc finger, FYVE domain containing 26	ZFYVE26	23503	NP_056161	NM_015346	4D	213073_at
nuclear protein UKp68	FLJ11806	79882	NP_079100	NM_024824	4D	213063_at
Nterminal asparagine amidase	NTANI	123803	NP_775745	NM_173474	4D	213062 at
microtubule associated serine/threonine kinase 3	MAST3	23031	XP_038150	XM_038150	4D	213045 at
ocular developmentassociated gene	ODAG	57798	NP_066990	NM_021167	4D	213018_at
tripartite motificontaining 37	TRIM37	4591	NP_056109	NM_015294	40	213009 s at
angiopoietinlike 2	ANGPTL2	23452	NP_036230	NM_012098	40	213001_at
KIAA0870 protein	KIAA0870	22898	NP 055772	NM_014957	4D	212974 at
ribose 5phosphate isomerase A (ribose 5phosphate e	RPIA	22934	NP_653164	NM_144563	ŧ	212973_at
					j	10070 04

hynothetical protein LOCOSASSO	LOC284352	284352	XP 113978	XM_113978	4D	213751 at
MAXlike protein X	MLX	6945	NP 937848	NM_198205	ŧ	213/08 S at
Rab geranylgeranyltransferase, beta subunit	RABGGTB	5876	NP_004573	NM 004582	t	21370° at
ribosomal protein L35a	RPL35A	6165	NP 000987	NM 000996	ŧ	21368/ s at
NOL1/NOP2/Sun domain family, member 5	NSUNS	55695	NP 683759	NM_148956	45	213670 x at
septin 6	SEPT6	23157	NP_665799	NM_145800	40	213666_at
zinc finger protein 75 (D8C6)	ZNF75	7626	NP_009062	NM_007131	4D	213659_at
similar to KIAA0363	LOC392666	392666	XP_374432	XM_374432	4D	213630 at
kinesin family member 3A	KIF3A	11127	NP_008985	NM_007054	4D	213623_at
NAD kinase	FLJ13052	65220	NP_075394	NM_023018	4D	213607 x at
FUS interacting protein (serinearginine rich)	FUSIPI	10772	NP_473357	NM_054016	4D	213594_x_at
angiotensin II receptorlike	AGTRL1	187	NP_005152	NM_005161	4D	213592 at
ribosomal protein L14	RPL14	9045	NP_003964	NM_003973	4D	213588 x at
EIA binding protein p300	EP300	2033	NP_001420	NM_001429	4D	213579 s_at
squalene epoxidase	SQLE	6713	NP_003120	NM_003129	4D	213577_at
sorting nexin 3	SNX3	8724	NP_003786	NM_003795	40	213545_x_at
major histocompatibility complex, class II. DP alo	HLADPAI	3113	NP_291032	NM_033554	40	213537_at
similar to hypothetical protein MGC13138	LOC146542	146542	NP_660314	NM_145271	40	213527_s_at
cyclin E1	CCNEI	898	NP_476530	NM_057182	40	213523 at
lamının, alpha 2 (merosin, congenital muscular dys	LAMA2	3900	INI_000+17	THE COURT	1	
coagulation factor II (thrombin) receptorlike 1	F2RL1	2000	NE 000417	TLTCOO MIN	4) (213519 s at
acylCoenzyme A oxidase 1, palmitoy1	ACOAL	2150	NID 005222	CVCS00 WIN	40	213506 at
collagen, type 11, alpha 1 (primary osteoarthritis	COLZAI	1200	2016+1 INI	NM 007202	40	13501 at
K1AA0073 protein	KIAA0073	23398	NP 036157	NM 022150	4) 6	213492 at
vesicleassociated membrane protein 4	VAMP4	8674	NP 973723	NM_201994	ŧ	213480 at
mangnant librous histiocytoma amplified sequence	MEHASI	7230	_004210			112400
Run an	To the state of th	350	SICKOO DIN	SCCOOL MN	40	213457 at
similar to RIIN and EVVE down	LOC441022	441022	XP 496700	XM 496700	4D	213437_at
enimorahin	EPIM	2054	NP 919337	NM 194356	4D	213434 at
ribosomal protein S19	RPS19	6223	NP_001013	NM_001022	4D	213414 s at
tight junction protein 3 (zona occludens 3)	TJP3	27134	NP_055243	NM_014428	4D	213412_at
hypothetical protein LOC 26208	TOC120208	120200	111 001002000			

NP_083051 80765 STARD5 START domain containing 5	NM_004315 NM_015033 NM_000992 NM_000993 NM_001003803 NM_0010038 NM_00100180 NM_00100180	at 4D	213902 at 213902 at 213903 x at 213909 x at 213909 x at 213905 at 213905 at 214002 at 214003 x at 214004 s at 214004 s at 214004 x at 214004 x at 214006 x at 2140
80765 STARDS	803		213902 at 213908 x 213908 x 213908 x 213908 at 213974 at 213995 at 214002 at 214003 x 214001 at 214001 at 214001 at 214000 at
80765 STARDS	803		213902 at 213909 x 213909 x 213909 at 213905 at 214002 at 214003 x 214011 s 214014 at 214040 s 214040 at 214040 at
80765 STARDS	803	at at at at	213902 at 213940 s 213940 s 213996 x 213995 at 214002 at 214013 x 214014 at 214014 s 214003 s 214014 s 214004 s
80765 STARDS	803	at at at at at	213902_at 213940_s 213969_x 213974_at 213995_at 214002_at 214003_x 214011_s 214014_at 214037_s
80765 STARDS	803	at at at	213902 at 213940 s 213969 x 213974 at 213995 at 214002 at 214003 x 214011 s 214014 at 214037 s
80765 STARDS	803	at at	213902 at 213940 s 213969 x 213974 at 213995 at 214002 at 214003 x 214011 s 214014 at
80765 STARDS	803	at at	213902_at 213940_s 213969_x 213974_at 213995_at 214002_at 214003_x 214011_s
80765 STARDS	803	at at	213902_at 213940_s 213969_x 213974_at 213995_at 214002_at 214003_x
80765 STARDS	803	at	213902_at 213940_s 213969_x 213974_at 213995_at 214002_at
80765 STARDS	803	at	213902_at 213940_s 213969_x 213974_at 213995_at
80765 STARDS		at	213902_at 213940_s 213969_x 213974_at
80765 STARDS		at	213902_at 213940_s 213969_x
80765 STARDS 30765 STARDS 40767 HOXAII 84705 GIPBP3 55062 WIPH9 1350 COX7C 1350 MGC52022 20526 ZCSI3 9487 PIGL 427 ASAHI 427 ASAHI 23048 FNBPI		at	213902_at
80765 STARDS			213902_at
80765 STARDS			
80765 STARDS		at 4D	213892 s at
80763 STARDS	NM_004278 NP_004269	4D	213889_at
80765 STARDS	NM_181706 NP_859057	4D	213853_at
80765 STARDS	NM_198563 NP_940965	4D	213851_at
80765 STARDS 9207 HOXA11 84705 GTPBP3 55062 WIPH9	NM_001867 NP_001858	4D	213846 at
80765 STARD5 3207 HOXA11 84705 GTPBP3	NM_017983 NP_060453	at 4U	213836_s_at
80765 STARD5 3207 HOXA11			213835 x at
80765 STARD5		4D	213823_at
		at 4D	213820 s
NP_003191 6929 TCF3 transcription factor 3 (E2A immunoglobulin enhance	NM_003200 NP_003191	at 4D	213811_x_at
NP_543031 5786 PTPRA protein tyrosine phosphatase, receptor type, A	NM_080841 NP_543031	at 4D	213799 s at
NP_006358 10487 CAP1 CAP, adenylate cyclascassociated protein 1 (yeast)	NM_006367 NP_006358	at 4D	213798 s
NP_006015 8887 TAX1BP1 Tax1 (human Tcell leukemia virus type I) binding	NM_006024 NP_006015	ŧ	213786_at
NP_963922 399563 FLJ43806 hypothetical protein FLJ43806	NM_201628 NP_963922	ŧ	213752_at

histone 1, H2bc	HIST IHZBC	834/	VICCOO AN	07CC00 IMIN	ŧ	214455_at
syntaxin 6	STX6	10228	NP 005810	NM 005819	ŧ	214441 at
Nacetyltransferase I (arylamine Nacetyltransferase	NATI	9	NP_000653	NM_000662	4D	214440_at
H2.0like homeo box 1 (Drosophila)	HLXI	3142	NP_068777	NM_021958	4D	214438_at
heat shock 70kDa protein 12A	HSPA12A	259217	XP_048898	XM_048898	4D	214434_at
nucleolar protein 1, 120kDa	NOLI	4839	NP_006161	NM_006170	4D	214427_at
methylCpG binding domain protein 2	MBD2	8932	NP_056647	NM_015832	4D	214397_at
arachidonate 5lipoxygenase	ALOX5	240	NP_000689	NM_000698	4D	214366_s_at
tropomyosin 3	TPM3	7170	NP_689476	NM_152263	4D	214365_at
acetylCoenzyme A carboxylase alpha	ACACA	31	NP_942131	NM_198834	4D	214358_at
decarboxy	0.000				i	
dona decarbovulare (aromatic) amino acid	חחר	1644	NP 000781	NM 000790	45	214347 s at
mitogenactivated protein kinase kinase kinase	MAP4K1	11184	NP 009112	NM_007181	4D	214339 s at
nephroblastoma overexpressed gene	NOV	4856	NP_002505	NM_002514	4D	214321_at
septin 6	SEPT6	23157	NP 055944	NM_015129	4D	214298 x at
ribosomal protein L17	RPL17	6139	NP_000976	NM_000985	4D	214291_at
fatty acid binding protein 3, muscle and heart (ma	FABP3	2170	NP_004093	NM_004102	4D	214285_at
galanin	GAL	51083	NP_057057	NM_015973	4D	214240_at
protein (peptidylprolyl cis/trans isomerase) NIMAi	PIN4	5303	NP_006214	NM_006223	4D	214224_s_at
complement component 1, q subcomponent binding pro	CIQBP	708	NP_001203	NM_001212	4D	214214_s_at
solute carrier family 25 (mitochondrial carrier; p	SLC25A17	10478	NP_006349	NM_006358	4D	214210_at
DiGeorge syndrome critical region gene 2	DGCR2	9993	NP_005128	NM_005137	4D	214198 s_at
tripeptidyl peptidase l	TPP1	1200	NP_000382	NM_000391	4D	214195_at
nuclear factor (erythroidderived 2)like 1	NFE2L1	4779	NP_003195	NM_003204	4D	214179_s_at
fumarate hydratase	FH	2271	NP_000134	NM_000143	4D	214170_x_at
succinate dehydrogenase complex, subunit B, iron s	SDHB	6390	NP_002991	NM_003000	4D	214166_at
polymerase (RNA) II (DNA directed) polypeptide D	POLR2D	5433	NP_004796	NM_004805	4D	214144_at
ribosomal protein L24	RPL24	6152	NP_000977	NM_000986	4D	214143 x at
nudix (nucleoside diphosphate linked moiety X)type	NUDT13	25961	NP_056985	NM_015901	4D	214136_at
pericentriolar material 1	PCM1	5108	NP_006188	NM_006197	4D	214118_x_at

Oprovem signaming moderator o (Accounte, C.	0.000	000000	01.0000	A TATAL CHIMNE S CO.	-	
Corotein cignalling modulator 3 (ACC213La C	GPSM3	07019	NP 071390	NM 022107	4D	214847 s at
docking protein 5	DOKS	55816	NP 060901	NM 018431	4D	214844 s at
hypothetical protein from clone 24828	KIAA1747	57795	NP_066988	NM_021165	4D	214822 at
exportin 6	XPO6	23214	NP_055986	NM_015171	4D	214784 x at
putative MAPK activating protein	MGC3794	261726	NP_690866	NM_152902	4D	214773 x at
G2 protein	G2	25758	XP_039515	XM_039515	4D	214772_at
Spl transcription factor	SPI	6667	NP 612482	NM_138473	4D	214732 at
zinc finger protein 200	ZNF200	7752	NP_003445	NM_003454	4D	214706_at
InaDlike (Drosophila)	INADL	10207	NP_795353	NM_176878	4D	214705_at
hypothetical protein MGC14376	MGC14376	84981	NP_001001870	NM_001001870	4D	214696_at
neuropilin 2	NRP2	8828	NP_958436	NM_201279	4D	214632_at
glucosidase, alpha; neutral AB	GANAB	23193	NP_938148	NM_198334	4D	214626_s_at
CASP8 and FADDlike apoptosis regulator	CFLAR	8837	NP_003870	NM_003879	4D	214618 at
tetraspan 2	TSPAN2	10100	NP_005716	NM_005725	4D	214606_at
ubiquitinconjugating enzyme E2D 1 (UBC4/5 homolog,	UBE2D1	7321	NP_003329	NM_003338	4D	214590_s_at
vacuolar protein sorting 52 (yeast)	VPS52	6293	NP 072047	NM_022553	4D	214585 s at
tumor necrosis factor receptor superfamily, member	TNFRSF21	27242	NP_055267	NM_014452	4b	214581_x_at
quaking homolog, KH domain RNA binding (mouse)	QKI	9444	NP_996735	NM_206853	4D	214543_x_at
quaking homolog, KH domain RNA binding (mouse)	QKI	9444	NP_996736	NM_206854	\$	214541_s_at
histone 1, H2bo	HIST1H2BO	8348	NP_003518	NM_003527	4D	214540 at
serine (or cysteine) proteinase inhibitor, clade B	SERPINB10	5273	NP_005015	NM_005024	4D	214539_at
				NM_005394	4D	214526 x at
mutL homolog 3 (E. coli)	MLH3	27030	NP_055196	NM_014381	4D	214525_x_at
Fc fragment of lgG, high affinity la, receptor (CD)	FCGRIA	2209	NP_000557	NM_000566	4D	214511_x_at
cAMP responsive element modulator	CREM	1390	NP_898883	NM_183060	4D	214508_x_at
exosome component 2	EXOSC2	23404	NP_055100	NM_014285	4D	214507_s_at
agouti signaling protein, nonagouti homolog (mouse	ASIP	434	NP_001663	NM_001672	4D	214498_at
sarcoglycan, delta (35kDa dystrophinassociated gly	SGCD	6444	NP_000328	NM_000337	4b	214492_at
serum amyloid A1	SAAI	6288	NP_000322	NM_000331	4D	214456_x_at

sect family domain containing 1 SMA4 RAB6 interacting protein 2 mevalonate kinase (mevalonic aciduria) postmetotic segregation increased 2like 5	PRIOZES po					
I family domain containing 1 A4 B6 interacting protein 2 valonate kinase (mevalonic aciduria)		5383	NP_777590	NM_174930		21566/ x at
1 family domain containing 1 A4 B6 interacting protein 2	MVK me		000422	NM_000431		215649 s at
1 family domain containing 1 A4	RAB6IP2 R/	5	-	NM_178040		215606 s at
1 family domain containing 1	SMA4 SN	11039	NP_067684	NM_021652	40	215599 at
	SCFD1 sec	23256	NP_057190	NM 016106		215548 s at
diaphanous homolog 1 (Drosophila)	DIAPH1 dia	1729	NP_005210	NM_005219		215541_s_at
butyrophilin, subfamily 2, member AI	BTN2A1 bu	11120	NP_008980	NM_007049		215493 x at
brain expressed Xlinkedlike 1	BEXL1 bra	56271	XP_043653	XM_043653		215440 s at
bromodomain adjacent to zinc finger domain, 2A	BAZZA br	11176	NP_038477	NM_013449	4D	215437 x at
troponin T2, cardiac	TNNT2 tro	7139	NP_001001430	NM_001001430	4D	215389 s_at
CD8 antigen, beta polypeptide 1 (p37)	CD8B1 CI	926	NP_742099	NM_172101	4D	215332 s_at
Rap2binding protein 9	RPIB9 Ra	154661	NP_612147	NM_138290	4D	215321_at
solute carrier family 12 (sodium/chloride transpor	SLC12A3 so	6559	NP_000330	NM_000339	4D	215274_at
transcription factor 3 (E2A immunoglobulin enhance	ICF3 tra	0929	NF_005191	NW 003200	ě	2 0 00 0 ut
eukaryotic translation initiation factor 3, subuni	000	8663		NM 003/52		215260 c at
keratin, hair, basic, 6 (monilethrix)	KRTHB6 ke	3892	NP_002275	NM 002284		215189 at
cadherinlike 22	CDH22 ca	64405	NP_067071	NM_021248	4	215181_at
NAD kinase	FLJ13052 N.	65220	NP_075394	NM_023018	4D	215159 s at
exosome component 8	EXOSC8 ex	11340	NP_852480	NM_181503	4D	215136_s_at
similar to K1AA0752 protein	LOC389347 sii	389347	XP_371783	XM_371783	4D	215133 s_at
KIAA0492 protein	K1AA0492 K	57238	XP_378914	XM_378914	4D	215109_at
NAD(P) dependent steroid dehydrogenaselike	NSDHL N.	50814	NP_057006	NM_015922	4D	215093 at
erythropoietin receptor	EPOR er	2057	NP_000112	NM_000121	4D	215054 at
neurexin 3	NRXN3 ne	9369	NP_004787	NM_004796	4D	215021_s_at
fasciculation and elongation protein zeta 2 (zygin	FEZ2 fa	9637	NP_005093	NM_005102	4D	215000 s_at
glutamate receptor, ionotropic, kainate 5	GRIK5 gl	2901	NP_002079	NM_002088	4D	214966 at
hypothetical protein MGC26885	MGC26885 hy	124044	NP_689552	NM_152339	4D	214965_at
KIAA0527 protein	KIAA0527 K	26032	XP_171054	XM_171054	40	214954 at
ATPase, Class II, type 9B	ATP9B A	374868	NP 940933	NM_198531	40	214934 at
SHC (Src homology 2 domain containing) transformin	SHC	Q Q	000020			
06			NP 003020	NM 003029	45	214853 s at

amid similar to dl.133P16.1 (ADPribosylation factor 1) superoxide dismutase 2, mitochondrial protein tyrosine phosphatase, nonreceptor type 12	protein	PIPNI2		NP 004072	NM 004081	4D	216922 x at
to dJ133P16.1 (ADPribosylation factor 1) ide dismutase 2, mitochondrial	or Location	C I V				-	
to dJ133P16.1 (ADPribosylation factor 1)	2		5782	NP 002826	NM_002835	4D	210915 s at
to dJ133P16.1 (ADPrihosylation factor 1)	superox	SOD2	6648	NP_000627		45	
4	similar	LOC442711	442711	XP_499457		ŧ	
glycine amidinotransferase (Larginine:glycine	glycine amid	GATM				45	
olfactory receptor, family 2, subfamily B, member	olfactor	OR2B6	2			45	216733 s at
prothymosin, alpha (gene sequence 28)	prothyn	PTMA			NM 012223	àŧ	216522 at
myeloid/lymphoid or mixedlineage leukemia (trithor	myeloid (trithor	MLLT10	8028		NW 00000	9 6	216515 v at
thymosinlike 3	thymos	TMSL3	7117		NM 004641	5 6	216503 s at
ATPbinding cassette, subfamily D (ALD), member 1	ATPbindi member l	ABCDI	215		NIM 102000	3 6	216438
leukotriene B4 receptor	leukotr	LTB4R	1241		NM 000027	t t	216418 at
)690	KIAA0690	KIAA0690	23223		NM 015179	ð	216388 ° at
deleted in azoospermia 4	deleted	DAZ4	57135	NP_065153	NM_020420	40	216360 x at
glycerol kinase	glycerc	GK	2710	NP 000158	NM_000167	ŧ	210316 X at
glutamate recentor metabotronic 8	glutam	GRM8	2918	NP_000836	NM_000845	4D	216255 s_at
parvin heta	parvin heta	PARVB	29780	NP_001003828	NM_001003828	4D	216254 at
Fas (TNE recentor subjunity 4, group A, member 2	Fac (T	FAS	355	NP_690612	NM_152873	4D	216252 x_at
nuclear recentor subfamily 4	nuclea	NR4A2	4929	NP_775263	NM_173171	4D	216248 s at
protein kinnse CAMPdenordom chain base subun	protein	PRKACA	5566	NP_002721	NM_002730	4D	216234_s_at
nalmitoultransferon	serine	SPTLC2	9517	NP_004854	NM_004863	4D	216202 s_at
KIRRA protein	KIBR	KIBRA	23286	NP_056053	NM_015238	4D	216074 x at
suppressor of hoir wine hard	Silvein	SUHWI	129025	NP_542778	NM_080740	4D	216034_at
shrine recent a lab.	olvoin	GLRA3	8001	NP 006520	NM_006529	4D	216021_s_at
ADB-ib-color o	A Date	ARERDI	10139	NP 003215	NM_003224	4D	215984_s_at
ominime decarboxylase antizyme [OHILI	D8S2298E	7993	NP 005662	NM_005671	4D	215983_s_at
hypothetical protein FLJ14668	riypou	OA71	4946	NP 004143	NM_004152	4D	215952 s_at
hacetylgiucosaminidase, alpha (Sanfilippo disease	Ivacet	FI 114668	84908	NP 116211	NM_032822	4D	215947 s at
ICK gamina alternate reading frame protein	ICK	NAGITI	4669	NP 000254	NM_000263	4D	215880_at
Poly (ADPribose) polymerase family, member 2	pory (TARP	445347	NP 001003806	NM_001003806	4D	215806_x_at
ras (1 Nr receptor superfamily, member 6)	ras (PARPS	85001	NP 005475	NM 005484	4D	215773_x_at
NE CONTRACTOR OF THE CONTRACTO	Enc /	FAS	355	NP 690613	NM_152874	4D	215719 x at

000				INM UIX/4/	ť	12 C C+/ / 17
preBcell colony enhancing factor 1	PBEF1	10135	NP 005/3/	NM 005/46	5	217742 S dt
preBcell colony enhancing factor 1	PBEFI	10135	NP_005737	NM 005/46	ŧ	217720 at
WD repeat domain 6	WDR6	11180	NP 060501	NM 018031	ŧ	217/34 S at
thymosin, beta 10	TMSB10	9168	NP_066926	NM 021103	6	217/35 s at
integral membrane protein 2B	ITM2B	9445	NP 068839	NM 021999	40	217/32 s at
S100 calcium binding protein A6 (calcyclin)	S100A6	6277	NP 055439	NM_014624	45	217728_at
solute carrier family 16 (monocarboxylic acid tran	SLC16A3	9123	NP_004198	NM_004207	4D	217691_x_at
winglesstype MMTV integration site family, member	WNI/B	/4//	NF_470079	1400 TAIN	t	- m
GTP binding protein 4	GTPBP4	23560	NP_036473	NM_012341	ŧ	217681 at
cation channel, sperm associated 2	CATSPER2	117155	NP_742095	NM_172097	40	217588 at
phenylalanine hydroxylase	PAH	5053	NP 000268	NM_000277	4D	217583_at
glutamate receptor, ionotrophic, AMPA 3	GRIA3	2892	NP_015564	NM_007325	4D	217565 at
				XM_497371	4D	217499 x at
phosphatase and tensin homolog (mutated in multipl	PTEN	5728	NP_000305	NM_000314	÷	21/492 s at
cytochrome c oxidase subunit VIIc	COX7C	1350	NP_001858	NM_001867	40	21/491 x at
interleukin 6 receptor	IL6R	3570	NP_000556	NM_000565	40	217489 s at
L1M domain kinase 2	LIMK2	3985	NP_005560	NM_005569	4D	217475 s_at
adrenergic, beta3, receptor	ADRB3	155	NP_000016	NM_000025	4D	217303_s_at
Nijmegen breakage syndrome 1 (nibrin)	NBSI	4683	NP_002476	NM_002485	4D	217299 s_at
mutL homolog 3 (E. coli)	MLH3	27030	NP_055196	NM_014381	40	217216 x at
KIAA1078 protein	KIAA1078	23271	NP 982284	NM_203459	4D	217196 s at
zinc finger protein 259	ZNF259	8882	NP_003895	NM_003904	4D	217186_at
glycerol kinase	GK	2710	NP_976325	NM_203391	4D	217167_x_at
mucin 4, tracheobronchial	MUC4	4585	NP_004523	NM_004532	4D	217109_at
itchy homolog E3 ubiquitin protein ligase (mouse)	ITCH	83737	NP_113671	NM_031483	Đ	217094_s_at
ELK1, member of ETS oncogene family	ELK1	2002	NP_005220	NM_005229	4D	217039 x at
hypothetical LOC389177	LOC389177	389177	XP_371680	XM_371680	4D	217016_x_at
alpha2glycoprotein 1, zinc	AZGPI	563	NP_001176	NM_001185	4D	217014 s at
RPA interacting protein	RIP	84268	NP_115684	NM_032308	4D	216962 at
Fc fragment of lgG, high affinity la, receptor (CD)	FCGRIA	2209	NP_000557	NM 000566	4D	216950 s at
retinoschisis (Xlinked, juvenile) 1	RSI	6247	NP_000321	NM_000330	4D	216937_s_at

	Clean	56851	NP 064539	NM 020154	4D	217898 at
NEFAinteracting nuclear protein NIP30	NIP30	80011	NP_079222	NM_024946	40	21 /896 s at
hypothetical protein FLJ13868	FLJ13868	64755	NP_073581	NM_022744	4	21/891 at
chromosome 2 open reading frame 25	C2orf25	27249	NP_056517	NM_015702	4	21/883 at
cell division cycle 27	CDC27	996	NP_001247	NM_001256	40	21/881 s_at
cell division cycle 27	CDC27	996	NP_001247	NM 001256	ŧ	2176/8 S at
macrophage migration inhibitory factor (glycosylat	MIF	4282	NP_002406	NM_002415	ŧ	21/8/1_s_at
Enah/Vasplike	EVL	51466	NP_057421	NM_016337	4	21/838 s at
vacuolar protein sorting 24 (yeast)	VPS24	51652	NP 057163	NM_016079	Đ	21/857 s_at
chromosome 20 open reading frame 24	C20orf24	55969	NP_061328	NM_018840	40	217835_x_at
synaptotagmin binding, cytoplasmic RNA interacting	SYNCRIP	10492	NP_006363	NM_006372	#	21 /834_s_at
NSFL1 (p97) cofactor (p47)	NSFLIC	55968	NP_057227	NM_016143	4	217831 s at
hypothetical protein FLJ13213	FLJ13213	79811	NP_079031	NM_024755	Đ	217828_at
actin related protein 2/3 complex, subunit 4, 20kD	ARPC4	10093	NP_005709	NM_005718	40	217818_s_at
suppressor of Ty 16 homolog (S. cerevisiae)	SUPT16H	11198	NP_009123	NM_007192	40	217815 at
selenoprotein T	SELT	51714	NP_057359	NM_016275	40	217811_at
Ufm1conjugating enzyme 1	Ufc1	51506	NP_057490	NM_016406	4	217797_at
RABIIB, member RAS oncogene family	RABIIB	9230	NP_004209	NM_004218	Đ	217793_at
G protein pathway suppressor	GPSI	2873	NP_004118	NM_004127	4D	217782 s_at
hypothetical protein HSPC152	HSPC152	51504	NP_057488	NM_016404	4Đ	217774 s at
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex	NDUFA4	4697	NP_002480	NM_002489	40	21///3_S_at
chromosome 13 open reading frame 12	C13orf12	51371	NP_057016	NM_015932	ŧ	217/69 s at
chromosome 14 open reading frame 166	C14orf166	51637	NP_057123	NM_016039	45	217768 at
small membrane protein !	SMPI	23585	NP_055128	NM_014313	45	217766_s_at
RAB31, member RAS oncogene family	RAB31	11031	NP_006859	NM 006868	40	217764_s_at
RAB31, member RAS oncogene family	RAB31	11031	NP_006859	NM_006868	40	217763 s at
membranetype I matrix metalloproteinase cytoplasmi	MTCBPI	55256	NP_060739	NM 018269	ŧ	217701_at
tripartite motifcontaining 44	TRIM44	54765	NP_060053	NM_017583	ŧ	217761 at
small EDRKrich factor 2	SERF2	10169	NP_005761	NM_005770	ŧ	217/56 x at
hematological and neurological expressed [HNI	51155	NP_057269	NM_016185	40	217755_at
adiponectin receptor 1	ADIPORI	21094	141 027002			

217902 s at	4b	NM 004667	NP 004658	8974	HERCO	hart domain and BI Da
217916 s_at	4D	NM_016623	NP_057707	51571	FAM49B	family with sequence similarity 49 member B
217918_at	4D	NM_014183	NP_054902	83658	DNCL2A	dynein, cytoplasmic, light polypeptide 2A
217926_at	4D	NM_014047	NP_054766	28974	HSPC023	HSPC023 protein
217927_at	4D	NM_014041	NP_054760	28972	SPCS1	signal peptidase complex subunit 1 homolog (S. cer
217940 s at	4D	NM_018210	NP_060680	55739	FLJ10769	hypothetical protein FLI10769
217949 s at	4D	NM 024006	NP_076869	79001	VKORC1	vitamin K epoxide reductase complex, subunit
217962_at	4D	NM_018648	NP_061118	55505	NOLA3	nucleolar protein family A, member 3 (11/ACA small
217971_at	4D	NM_021970	NP 068805	8649	MAP2K11P1	mitogenactivated protein kinase kinase interacti
217977_at	4D	NM_016332	NP 057416	51734	SEPXI	selenoprotein X. 1
217988 at	4D	NM_021178	NP_067001	57820	CCNB11P1	cyclin B1 interacting protein
217994 x at	4D	NM_017871	NP 060341	54973	FLJ20542	hypothetical protein FLI20542
217995_at	4D	NM_021199	NP_067022	58472	SQRDL	sulfide quinone reductaselike (veast)
218001_at	4D	NM_016034	NP_057118	51116	MRPS2	mitochondrial ribosomal protein \$2
218005_at	4D	NM_006963	NP_008894	7570	ZNF22	zinc finger protein 22 (KOX 15)
218007 s at	4D	NM_015920	NP_057004	51065	RPS27L	ribosomal protein S27like
218010 x at	4D	NM_024299	NP_077275	79144	C20orf149	chromosome 20 open reading frame 149
218022_at	4D	NM_016440	NP_057524	51231	VRK3	vaccinia related kinase 3
218032_at	4D	NM_003498	NP_003489	8303	SNN	stannin
218033 s at	45	NM_003498	NP_003489	8303	SNN	stannin
218049 s at	4Đ	NM_014078	NP 054797	28998	MRPL13	mitochondrial ribosomal protein 1.13
218056_at	4D	NM_016561	NP_057645	51283	BFAR	bifunctional apoptosis regulator
218059_at	4D	NM_016096	NP_057180	51123	LOC51123	HSPC038 protein
218061_at	4D	NM_014623	NP_055438	4201	MEA	maleenhanced antigen
218073_s_at	4D	NM_018087	NP_060557	55706	FLJ10407	hypothetical protein FLJ 10407
218074 at	4D	NM_016062	NP 057146	51647	CG1128	CG1128 protein
218077 s_at	4D	NM 016598	NP_057682	51304	ZDHHC3	zinc finger, DHHC domain containing 3
218081 at	4D	NM_017874	NP_060344	54976	C20orf27	chromosome 20 open reading frame 27
218089 at	4D	NM_015511	NP_056326	25980	C20orf4	chromosome 20 open reading frame 4
218094 s at	40	NM_018478	NP 060948	55861	C20orf35	chromosome 20 open reading frame 35
210090_at	ŧ	NAI 018361	NP_060831	55326	AGPAT5	lacylglycerol3phosphate Oacyltransferase 5 (lysoph

The state of the s	0058486	58486	NP 06/034	INM 021211	ŧ	710702 S SI
polymerase (RNA) I polypeptide D, 16kDa	POLRID	28010	05/056	NIM 033312	5 6	10 00201
fibrosin I	FBSI	64319	NP_071897	NM 0122452	4 5	218258 at
zinc finger, DHHC domain containing 6	ZDHHC6	64429	NP_071939	NM_022494	ŧ	
nucleolar protein 8	NOL8	55035	NP 060418	NM_017948	4D	218244 at
				NM_017601	4D	218233 s at
pogo transposable element with KRAB domain	POGK	57645	NP 060012	NM_017542	40	218229 S at
NADH dehydrogenase (ubiquinone) I beta subcomplex,	NDUFB4	4710	NP_004538	NM_00454/	ŧ	210220 S at
LanC lantibiotic synthetase component Clike 2 (bac	LANCL2	33913	001107	Nit 0010071	7	210226
chromosome 11 open reading frame 10	Cl lorf10	746	NP 055021	NM 019607	5 8	218213 s at
fructosamine3kinaserelated protein	FN3KRP	79672	NP 078895	NM_024619	ŧ	218212 at
nucleolar protein family 6 (RNAassociated)	NOL6	65083	NP_075068	NM_022917	ŧ	218210 S at
osteopetrosis associated transmembrane protein 1	OSTMI	28962	NP_054747	NM_014028	4D	218196_at
ubiquinolcytochrome c reductase complex (7.2 kD)	HSPC051	29796	NP_037519	NM_01338/	ŧ	210190_S_at
vacuolar protein sorting 4B (yeast)	VPS4B	9525	NP_004860	NM_004869	ŧ	2181/1 at
ADPribosylation factorlike 5	ARLS	26225	NP_036229	NM_012097	4D	218150 at
signal recognition particle receptor, B subunit	SRPRB	58477	NP_067026	NM_021203	4D	218140 x at
NIF3 NGG1 interacting factor 3like 1 (S. pombe)	NIF3L1	60491	NP_068596	NM_021824	40	218133 s at
leukocyte receptor cluster (LRC) member 5	LENG5	79042	NP_076980	NM_024075	4D	218132_s_at
chromosome 21 open reading frame 59	C21orf59	56683	NP_060305	NM_017835	4D	218123_at
translocase of inner mitochondrial membrane 23 hom	TIMM23	10431	NP_006318	NM_006327	ŧ	710110_2_at
ringbox 1	RBX1	9978	NP 055063	NM 014248	ŧ	21011/ at
chromosome 9 open reading frame 78	C9orf78	51759	NP 057604	NM 016520	ŧ	218116_at
ASF1 antisilencing function 1 homolog B (S. cerevi	ASFIB	55723	NP_060624	NW_018154	ŧ	TID 11013
testis expressed sequence 10	TEX10	54881	NP_060216	NM_017746	ŧ	218104 at
NADH dehydrogenase (ubiquinone) 1, subcomplex unkn	NDUFC2	4/18	NF_004540	NW 004349	ŧ	3 01013
uncharacterized hypothalamus protein HT008	HT008	55852	NP 060939	NM 018469	ŧ	218101 a at
ADPribosylation factor guanine nucleotideexchange	ANTOEF2	1000	NF_000411	MINI 000420	ė	210000

sorting nexm 10	SNXIO	19867	PC+/CO TNI	INIM 013322	10	- TOTOL W
miscr protein 201	CATALO	20007	ND OTAGA	CCCCIO MIN	Đ.	TIRADA at
zinc finger protein 281	ZNF281	23528	NP 036614	NM 012482	4D	218401 s at
RNA binding motif protein 7	RBM7	10179	NP_057174	NM_016090	4D	218379 at
PRKR interacting protein 1 (IL11 inducible)	PRKRIPI	79706	NP_078929	NM_024653	40	2185/8 s at
nudix (nucleoside diphosphate linked moiety X)type	NUDT9	53343	NP_076952	NM_024047		2185/5_at
chromosome 12 open reading frame 4	C12orf4	57102	NP_065107	NM_020374		2183/4 s at
FLJ20859 gene	FLJ20859	64745	NP_073571	NM_022734		218366_x_at
chromosome 14 open reading frame 114	C14orf114	55218	NP_060669	NM_018199	4D	218363_at
translocase of inner nutochondrial membrane 8 homo	HIMMINI	20321	INI_020271	0.2455	t	
FtsJ homolog 2 (E. coli)	FTSJ2	29960	NP_037525	NM_013393	ŧ	219357 6 01
Zwilch	FLJ10036	55055	NP_060445	NM_017975	10	218349 S at
retinoic acid induced 16	RAI16	64760	NP_073586	NM_022749	4D	218337_at
prefoldin 2	PFDN2	5202	NP_036526	NM_012394	40	218336_at
brain expressed, Xlinked I	BEX1	55859	NP 060946	NM_018476	4D	218332_at
coenzyme Q4 homolog (yeast)	COQ4	51117	NP_057119	NM_016035	4D	218328_at
spermatogenesis associated, serinerich 2	SPATS2	65244	NP_075559	NM_023071	4D	218324 s at
pellino homolog 1 (Drosophila)	PELII	57162	NP_065702	NM_020651	4D	218319_at
translocase of inner mitochondrial membrane 9 homo	MMI	20020	NF_030392	1012400	ŧ	
CDK5 regulatory subunit associated protein 1	CDKSKAPI	21024	NF 03/492	004010 MIN	à	218316 of
calcium/calmodulindependent protein kinase II	CaMKIINalpha	53654	NF 001034	NM 016408	45	218315 s at
		20120	NID 061064	NM 019597	45	218309 at
chromosome 10 open reading frame 97	C10orf97	80013	NP 0/9224	NM 024948	ŧ	218207 at
nucleoporin 50kDa	NUP50	10762	NP 009103	NM 00/1/2	ŧ	210293 S at
protein kinase, AMPactivated, gamma 2 noncatalytic	PRKAG2	51422	NP_057287	NM_016203	ŧ ė	218292 S at
mitogenactivated proteinbinding proteininteracting	MAPBPIP	28956	NP_054736	NM_014017	40	218291 at
ring finger protein 7	RNF7	9616	NP_055060	NM_014245	4D	218286 s at
dehydrogenase/reductase (SDR family) member 6	DHRS6	56898	NP_064524	NM_020139	4D	218285 s_at
synovial sarcoma translocation gene on chromosome	SS18L2	51188	NP_057389	NM_016305	4	218285_at
mitochondrial ribosomal protein L24	MRPL24	79590	NP_078816	NM 024540	40	2182/0_at
BRCA2 and CDKN1A interacting protein	BCCIP	56647	NP_057651	NM_016567	4D	218264_at

	NP_037498	NM_013366	ŧ	
		220010 1111	ָרָיִי קיי	218555_at
	NP 060751	NM_018281	4D	218552_at
57380 MRS2L	NP_065713	NM_020662	4 D	218538_s_at
80194 FLJ21749	NP_079400	NM_025124	40	218531_at
54840 APTX	NP_060162	NM_017692	4D	218527_at
		NM_014185	40	218526_s_at
94104 C21orf66	NP_057715	NM_016631	4D	218515_at
55181 FL	NP_060619	NM_018149	4D	218514_at
55759 W	NP_060726	NM_018256	4D	218512 at
7227 TF	NP_054831	NM_014112	4D	218502 s_at
8409 UX	NP 004173	NM_004182	4D	218495_at
55900 ZN	NP_060913	NM_018443	4D	218490_s_at
56912 FL	NP_064538	NM_020153	4D	218483_s_at
56915 EX	NP_064543	NM_020158	4D	218481_at
582 BI	NP_078925	NM_024649	4D	218471_s_at
51067 CC	NP_057020	NM_015936	4D	218470_at
	1			
	NP 064617	CECOCO MN	4	218467 at
	NP 079404	NM 025128	4D	218463 s at
	NP 057385	NM 016301	4D	218461 at
54919 FL	NP_060272	NM_017802	4D	218460_at
9054 NI	NP_066923	NM_021100	4D	218455_at
79887 FL	NP_079105	NM_024829	4D	218454_at
64866 CI	NP 073753	NM_022842	4D	218451_at
55325 FL	NP_060829	NM_018359	4D	218449_at
56942 DO	NP_064573	NM_020188	4D	218447_at
26015 RF	NP_056355	NM_015540	4D	218441_s_at
56922 M	NP_064551	NM_020166	ŧ	218440_at
80306 M	NP_079481	NM_025205	4D	218438_s_at
10807 SI	NP_006634	NM_006643	4D	218427_at
PENDIX D	AP			
מייאומוגיים ה	À			
	PENDIX D 10807 SDCCAG3 80306 MED28 80306 MED28 80922 MCCC1 26042 DC13 50942 DC13 50943 FLJ2062 50964 NES1 50964 MUS81 5097 CC104 5098 MUS8 5097 CC104 5097 FLJ21827 5099 UXT 5099 UXT	APPENDIX D 10807 80306 80306 80306 26015 26015 250922 26015 36942 350942 350943 31184 31067 382 36912 36912 350912 350912 350912 35739	APPENDIX D NP_006634 10807 NP_079481 80336 NP_064551 56922 NP_065355 560415 NP_064573 56942 NP_066773 56947 NP_079105 79887 NP_079105 79887 NP_079105 79887 NP_079105 51867 NP_079404 80198 NP_064617 56984 NP_079404 80198 NP_064538 56915 NP_064538 56912 NP_064538 56912 NP_064538 56902 NP_064581 55900 NP_064581 7227 NP_0646775 54878 NP_064173 54909 NP_064581 7227 NP_0660725 55789 NP_0660725 55789	NM_006643 NP_06634 10807 NM_025205 NP_064551 56922 NM_020166 NP_064551 56922 NM_015540 NP_064551 56942 NM_01888 NP_064553 56942 NM_020188 NP_066923 9054 NM_020829 NP_067925 96547 NM_021100 NP_067923 9054 NM_021100 NP_067923 9054 NM_021100 NP_067923 9054 NM_021100 NP_067923 9054 NM_02169 NP_067923 9054 NM_01630 NP_067923 9054 NM_01630 NP_067923 90592 NM_02163 NP_064517 56984 NM_02163 NP_064518 56912 NM_02163 NP_064518 56912 NM_018413 NP_064538 56912 NM_018412 NP_064726 55759 NM_018429 NP_064677 56981

RAB7, member RAS oncogene familylike 1	RAB7L1	8934	NP_003920	NM_003929	4D	218699 at
leucine rich repeat containing 5	LRRC5	55144	NP_060573	NM_018103	4D	218684_at
hypothetical protein FLJ13611	FLJ13611	80006	NP_079217	NM_024941	₽ P	218674 at
ATPase inhibitory factor 1	ATPIFI	93974	NP_057395	NM_016311	4D	218671_s_at
RAP2C, member of RAS oncogene family	RAP2C	57826	NP_067006	NM_021183	4D	218669 at
hypothetical protein FLJ14154	FLJ14154	79903	NP_079121	NM_024845	4D	218661_at
mitochondrial ribosomal protein S33	MRPS33	51650	NP 057155	NM_016071	4D	218654 s at
				NM_022775	4D	218650_at
transducer of regulated cAMP response elementbindi	TORC3	64784	NP_073606	NM_022769	4D	218648_at
hypothetical protein FLJ20534	FLJ20534	54969	NP_060337	NM_017867	4D	218646 at
postsynaptic protein CRIPT	CRIPT	9419	NP_054890	NM_014171	4D	218643 s at
coiledcoilhelixcoiledcoilhelix domain containing 7	CHCHD7	79145	NP_077276	NM_024300	4D	218642 s_at
				NM_023941	4D	218641_at
abhydrolase domain containing 10	ABHD10	55347	NP_060864	NM_018394	4D	218633 x at
arginine vasopressininduced 1	AVPII	60370	NP_068378	NM_021732	4D	218631 at
smoothened homolog (Drosophila)	SMO	6608	NP_005622	NM_005631	4D	218629_at
hypothetical protein FLJ11259	FLJ11259	55332	NP_060840	NM_018370	4D	218627_at
nudix (nucleoside diphosphate linked moiety X)type	NUDT2	318	NP_001152	NM_001161	4D	218609_s_at
SDA1 domain containing 1	SDADI	55153	NP_060585	NM_018115	4D	218607 s_at
transcription factor B2, mitochondrial	TFB2M	64216	NP_071761	NM_022366	4D	218605_at
LEM domain containing 3	LEMD3	23592	NP_055134	NM_014319	4D	218604_at
upregulated gene 4	URG4	55665	NP_060390	NM_017920	4D	218601_at
				NM_018072	4D	218594 at
progressive external ophthalmoplegia 1	PEO1	56652	NP_068602	NM_021830	4D	218590_at
chromosome 5 open reading frame 3	C5orf3	10827	NP_061161	NM_018691	4D	218588 s at
abhydrolase domain containing 4	ABHD4	63874	NP_071343	NM_022060	4b	218581_at
DEAH (AspGluAlaHis) box polypeptide 35	DHX35	60625	NP 068750	NM_021931	4D	218579 s at
hyperparathyroidism 2 (with jaw tumor)	HRPT2	79577	NP 078805	NM_024529	4D	218578_at
anaphase promoting complex subunit !	ANAPCI	64682	NP_073153	NM_022662	4D	218575 at
chromosome 14 open reading frame 123	C14orf123	29082	NP_054888	NM_014169	4D	218571_s_at
hypothetical protein FLJ10747	FLJ10747	55219	NP_060672	NM_018202	4D	218562 s_at
chromosome 6 open reading frame 149	C6orf149	57128	NP_065141	NM_020408	4D	218561 s at

chitobiase, diNacetyl	CTBS	1486	NP 004379	NM_004388	4D	218924 s_at
hypothetical protein FLJ14007	FLJ14007	79752	NP_078975	NM_024699	4D	218919_at
hypothetical protein FLJ10375	FLJ10375	55129	NP_060545	NM_018075	4D	218910 at
hypothetical protein FLJ20530	FLJ20530	55656	NP_060334	NM 017864	4D	218905 at
mitochondrial ribosomal protein L35	MRPL35	51318	NP_057706	NM_016622	4D	218890 x_at
MLF1 interacting protein	MLFIIP	79682	NP_078905	NM_024629	4D	218883_s_at
WD repeat domain 3	WDR3	10885	NP 006775	NM_006784	4D	218882_s_at
FOSlike antigen 2	FOSL2	2355	NP 005244	NM_005253	4D	218880_at
hypothetical protein FLJ12998	FLJ12998	64779	NP_073601	NM_022764	4D	218879 s at
hypothetical protein FLJ22390	FLJ22390	64757	NP_073583	NM_022746	4D	218865_at
WD repeat domain 33	WDR33	55339	NP_060853	NM_018383	4D	218851_s_at
hypothetical protein FLJ12788	FLJ12788	64427	NP_071937	NM_022492	4D	218838_s_at
phospholipid scramblase 3	PLSCR3	57048	NP_065093	NM_020360	4D	218828_at
aminopeptidaselike l	NPEPLI	79716	NP_078939	NM_024663	4D	218822_s_at
				NM_025156	4D	218812_s_at
chromosome 7 open reading frame 19	C7orf19	80228	NP_116220	NM_032831	4D	218811_at
vav 3 oncogene	VAV3	10451	NP_006104	NM_006113	4D	218807_at
thioredoxinlike 4B	TXNL4B	54957	NP_060323	NM_017853	4D	218794 s at
sex comb on midleglike 1 (Drosophila)	SCMLI	6322	NP_006737	NM_006746	4D	218793 s at
methionine sulfoxide reductase B2	MSRB2	22921	NP_036360	NM_012228	4D	218773_s_at
transmembrane protein 39B	ТМЕМЗ9В	55116	NP_060526	NM_018056	4D	218770_s_at
				NM_015940	4D	218760_at
hypothetical protein FLJ23323	FLJ23323	79707	NP_078930	NM 024654	4D	218754_at
hypothetical protein FLJ10307	FLJ10307	55113	NP_060523	NM_018053	4D	218753_at
hypothetical protein FLJ20422	FLJ20422	54929	NP_060284	NM 017814	4D	218745 x_at
chromosome 22 open reading frame 18	C22orf18	79019	NP_076958	NM_024053	4D	218741_at
ring finger protein 138	RNF138	51444	NP_057355	NM_016271	4D	218738_s_at
zinc finger protein 544	ZNF544	27300	NP_055295	NM_014480	4D	218735_s_at
latexin	LXN	56925	NP_064554	NM_020169	4D	218729 at
hypothetical protein FLJ12436	FLJ12436	79714	NP 078937	NM_024661	4D	218722 s_at
leprecanlike 1	LEPRELI	55214	NP_060662	NM_018192	4D	218717 s at
hepatocellular carcinomaassociated antigen 66	HCA66	55813	NP_060898	NM_018428	4D	218715_at
NTF2like export factor 1	ITXN	29107	NP_037380	NM_013248	4D	218708_at
lactamase, beta 2	LACTB2	51110	NP_057111	NM_016027	4D	218701_at

218926_at	4D	NM_018657	NP_061127	55892	NNYM	myoneurin
218939_at	4D	NM_012318	NP_036450	3954	LETMI	leucine zipperEFhand containing transmembrane
218942_at	4D	NM_024779	NP 079055	79837	PIP5K2C	phosphatidylinositol4phosphate 5kinase type II o
218949_s_at	4D	NM_018292	NP_060762	55278	QRSL1	glutaminyltRNA synthase
218953_s_at	4D	NM_024028	NP_076933	78991	MGC3265	hypothetical protein MGC3265
218968 s at	Đ	NM_018197	NP_060667	55734	ZFP64	zinc finger protein 64 homolog (mouse)
218978_s_at	4D	NM_018586				0 , 0 ,
218981_at	4D	NM_020186	NP 064571	57001	ACN9	ACN9 homolog (S. cerevisiae)
218982 s at	4D	NM_015969	NP_057053	51373	MRPS17	mitochondrial ribosomal protein S17
218984_at	4D	NM_019042	NP_061915	54517	FLJ20485	hypothetical protein FLJ20485
218986_s_at	4D	NM_017631	NP_060101	55601	FLJ20035	hypothetical protein FLJ20035
218993_at	4D	NM_018146	NP_060616	55178	RNMTLI	RNA methyltransferase like 1
218999 at	4D	NM_018295	NP_060765	55281	FLJ11000	hypothetical protein FLJ11000
219006_at	4D	NM_014165	NP_054884	29078	C6orf66	chromosome 6 open reading frame 66
219007_at	4D	NM_024647	NP_078923	348995	NUP43	nucleoporin 43kDa
219024_at	4D	NM_021622	NP_067635	59338	PLEKHAI	pleckstrin homology domain containing, family A
219027_s_at	4D	NM_006901	NP_008832	4649	MYO9A	myosin IXA
219030_at	4D	NM_016058	NP_057142	51002	CG1121	CG1121 protein
219032_x_at	4D	NM_014322	NP_055137	23596	OPN3	opsin 3 (encephalopsin, panopsin)
219033 at	4D	NM_024615	NP_078891	79668	PARP8	poly (ADPribose) polymerase family, member 8
219034 at	4D	NM_017851	NP_060321	54956	PARP16	poly (ADPribose) polymerase family, member 16
219037_at	4D	NM_016052	NP_057136	51018	CGII15	CG1115 protein
219040_at	4D	NM_024535	NP_078811	79585	CORO7	coronin 7
219053 s at	4D	NM 017966	NP_060436	55048	VPS37C	vacuolar protein sorting 37C (yeast)
219054_at	4D	NM_024563	NP_078839	79614	FLJ14054	hypothetical protein FLJ14054
219056_at	4D	NM_024570	NP_078846	79621	FLJ11712	hypothetical protein FLJ11712
219066_at	4D	NM_021823	NP_068595	60490	MDS018	hypothetical protein MDS018
219067 s at	4D	NM_017615	NP_060085	54780	C10orf86	chromosome 10 open reading frame 86
219069_at	4D	NM_017704	NP_060174	54851	FGIF	fetal globininducing factor
219079_at	4D	NM_016230	NP_057314	51167	NCB5OR	NADPH cytochrome B5 oxidoreductase
219093 at	4D	NM_017933	NP_060403	55022	FLJ20701	hypothetical protein FLJ20701

centrosome protein Cep63	cepos	40700	INI_0/2420	1001000 ININ		
	62.63	1200	ND 070456	08180 MN		219242 at
secemin 3	SCRN3	79634	NP 078859	NM 024583	4D	219234 x at
gasderminlike	GSDML	55876	NP_061000	NM 018530	4D	219233 s_at
zinc finger protein 331	ZNF331	55422	NP_061025	NM_018555	4D	219228_at
mitochondrial ribosomal protein S22	MRPS22	56945	NP_064576	NM_020191	4D	219220 x at
RAB8B, member RAS oncogene family	RAB8B	51762	NP 057614	NM_016530		219210 s_at
interferon induced with helicase C domain 1	IFIH1	64135	NP 071451	NM_022168		219209 at
ubiquitin associated protein 2	UBAP2	55833	NP_060919	NM_018449	4D	219192_at
eukaryotic translation initiation factor 2C, 4	EIF2C4	192670	NP_060099	NM_017629	4D	219190_s_at
pleckstrin homology, Sec7 and coiledcoil domains	PSCD4	27128	NP_037517	NM_013385	4D	219183_s_at
hypothetical protein FLJ22555	FLJ22555	79568	NP_078796	NM_024520		219176 at
ubiquitin domain containing 1	UBTDI	80019	NP_079230	NM_024954		219172_at
zinc finger protein 236	ZNF236	7776	NP_031371	NM_007345	4D	219171_s_at
mitochondrial ribosomal protein L11	MRPL11	65003	NP_057134	NM_016050	4D	219162_s_at
kelchlike 2, Mayven (Drosophila)	KLHL2	11275	NP_009177	NM_007246	4D	219157_at
synaptojanin 2 binding protein	SYNJ2BP	55333	NP_060843	NM_018373	4D	219156_at
podocalyxinlike 2	PODXL2	50512	NP_056535	NM_015720		219152_at
RAB, member of RAS oncogene familylike 2B	RABL2B	11158	NP_009012	NM_007081		219151 s at
debranching enzyme homolog 1 (S. cerevisiae)	DBRI	51163	NP_057300	NM_016216		219149 x at
hypothetical protein FLJ22729	FLJ22729	79736	NP_078959	NM_024683	4D	219146_at
hypothetical protein FLJ20604	FLJ20604	54995	NP_060367	NM_017897	4D	219133_at
hypothetical protein FLJ10287	FLJ10287	54482	NP_061956	NM_019083		219130_at
hypothetical protein FLJ20558	FLJ20558	54980	NP_060350	NM_017880	4D	219128_at
hypothetical protein MGC11242	MGC11242	79170	NP_077296	NM_024320	4D	219127_at
zinc finger protein 232	ZNF232	7775	NP_055334	NM_014519	4D	219123_at
hypothetical protein FLJ20171	FLJ20171	54845	NP_060167	NM_017697	4D	219121 s_at
LSM8 homolog, U6 small nuclear RNA associated (S.	LSM8	51691	NP_057284	NM_016200	40	219119_at
DEAD (AspGluAlaAsp) box polypeptide 54	DDX54	79039	NP 076977	NM_024072		219111 s at
DEAD (AspGluAlaAsp) box polypeptide 27	DDX27	55661	NP_060365	NM_017895		219108_x_at
ring finger protein 141	RNF141	50862	NP_057506	NM_016422	4D	219104_at
chromosome 12 open reading frame 5	C12orf5	57103	NP_065108	NM 020375	4D	219099_at
armadillo repeat containing 7	ARMC7	79637	NP_078861	NM_024585	4D	219096 at

NYI-4308244vI

219459_at 219467 at 219451 at 219442_at 219436 s at 219434 at 219419_at 219410_at 219398_at 219318_x_at 219397_at 219377 at 219373_at 219363 s at 219356_s_at 219345 at 219336 s at 219330_at 219329 s_at 219305_x_at 219292_at 219279 at 219276 x at 219269 at 219266_at 219244 s at 219243 at 219378_at 219367 s at 219312_s_at 219248 at 40 40 4D 4D 4D 4D 40 45 40 40 40 4 4 4 40 4D **4 4** 4 8 4 4D 4 40 4D 4D 4 NM_017676 NM_018082 NM_016064 NM_022094 NM_024561 NM_018973 NM_018534 NM_015942 NM_016074 NM_015947 NM 024062 NM_016085 NM_016060 NM_024048 NM 016242 NM_018643 NM_024805 NM_018004 NM_025147 NM_022751 NM_016410 NM_023929 NM_012168 NM_025264 NM 018326 NM_018105 NM_017718 NM_024828 NM_024567 NM 021632 NM_022163 NP_060146 NP_076953 NP_057326 NP 079081 NP 060474 NP 060796 NP_060552 NP_061113 NP_071377 NP 079423 NP_078837 NP_073588 NP 061846 NP_057494 NP_057158 NP_057031 NP_057144 NP_076418 NP_036300 NP 079104 NP_078843 NP_067645 NP_079540 NP_071446 NP_061004 NP_057026 NP_057169 NP_060575 54826 55703 51705 51374 5100 65986 54210 55076 63924 80219 79612 64762 54344 8828 51001 51510 51027 51008 26232 79863 55145 79886 79618 59348 80745 26589 55303 FLJ20125 C2orf28 POLR3B MGC3020 C18orf22 NARGIL NRP2 CG112 THAP TREMI TMEM45A CIDEC FLJ13448 C18orf11 DPM3 SNF7DC2 CG1143 ASCCI MED31 ZBTB10 FBXO2 MRPL46 GIMAP4 EMCN C9orf82 FLJ21616 ZNF350 THUMPD2 hypothetical protein FLJ20125 polymerase (RNA) III (DNA directed) polypeptid hypothetical protein MGC3020 triggering receptor expressed on myeloid cells 1 chromosome 18 open reading frame 22 transmembrane protein 45A NMDA receptor regulated 1 like SNF7 domain containing 2 CG1143 protein chromosome 2 open reading frame 28 mediator of RNA polymerase II transcription zinc finger and BTB domain containing 10 Fbox protein 2 cell deathinducing DFFAlike effector c hypothetical protein FLJ13448 chromosome 18 open reading frame polypeptide dolichylphosphate mannosyltransferase neuropilin 2 CG112 protein activating signal cointegrator 1 complex subunit chromosome 9 open reading frame 82 hypothetical protein FLJ21616 GTPase, IMAP family member 4 THAP domain containing, apoptosis associated zinc finger protein 350 THUMP domain containing 2 mitochondrial ribosomal protein 1.46

78

	FI 120280	54876	NP 060211	NM 017741	4D	219717 at
plexin domain containing 1	PLXDC1	57125	NP_065138	NM_020405	4D	219700_at
hypothetical protein FLJ22573	FLJ22573	79713	NP_078936	NM_024660	4D	219690_at
RAB11 family interacting protein 1 (class I)	RABITFIPI	80223	NP_079427	NM_025151	4D	219681_s_at
Kruppellike factor 3 (basic)	KLF3	51274	NP 057615	NM_016531	4D	219657 s at
protocadherin 12	PCDH12	51294	NP_057664	NM_016580	4D	219656 at
				NM_014054	4D	219653_at
developmental pluripotency associated 4	DPPA4	55211	NP_060659	NM_018189	4D	219651_at
popeye domain containing 2	POPDC2	64091	NP_071418	NM_022135	4D	219647 at
NYREN58 antigen	NYREN58	51134	NP_057206	NM_016122	4D	219644 at
deetiolated homolog 1 (Arabidopsis)	DET1	55070	NP_060466	NM_017996	4D	219641_at
				NM_024937	4D	219631_at
sirtuin (silent mating type information regulation	SIRT6	51548	NP_057623	NM_016539	4D	219613 s at
hypothetical protein DKFZp434L0117	DKFZP434L011	64793	NP_073615	NM_022778	4D	219611_s_at
membranespanning 4domains, subfamily A, member 4	MS4A4A	51338	NP_076926	NM_024021	Đ	219607_s_at
RWD domain containing I	RWDDI	51389	NP_057188	NM_016104	4D	219598 s at
ninjurin 2	NINJ2	4815	NP_057617	NM_016533	4D	219594_at
CG130 protein	CG130	51611	NP_057042	NM_015958	40	219590 x at
likely homolog of yeast SEN2	SEN2L	80746	NP_079541	NM_025265	Đ	219581_at
reticulon 3	RTN3	10313	NP_006045	NM_006054	4D	219549_s_at
FLJ22624 protein	FLJ22624	79866	NP_079084	NM_024808	40	219544 at
zinc finger protein 267	ZNF267	10308	NP_003405	NM_003414	4D	219540_at
chromosome 14 open reading frame 169	C14orf169	79697	NP_078920	NM_024644	4D	219526_at
transient receptor potential cation channel, subfa	TRPV4	59341	NP_067638	NM_021625	4D	219516_at
hypothetical protein FLJ13150	FLJ13150	79871	NP_079089	NM_024813	4D	219504_s_at
transmembrane protein 40	TMEM40	55287	NP_060776	NM_018306	4D	219503 s_at
Bcell CLL/lymphoma 11A (zinc finger protein)	BCL11A	53335	NP_060484	NM_018014	45	219498 s at
cysteinerich hydrophobic domain 2	CHIC2	26511	NP_036242	NM_012110	4D	219492_at
hypothetical protein FLJ23560	FLJ23560	79738	NP_078961	NM_024685	4D	219487_at
chromosome 21 open reading frame	C21orf18	54093	NP_059134	NM_017438	4D	219482_at
				NM_024115	40	219476_at
Building induced difficultiation associated blo	CONT	77077	141 OCT DO	1414 017000	-	21 /4/ J at

Ctype lectin domain family 4, member A	CLEC4A	50856	NP_057268	NM_016184	4D	219947_at
XX S	G	51022	NP 057150	NM 016066	4D t	219933 at
3 6	FGD6	2054	NP 060821	NM 021070	45	219901_at
35A	CLEC5A	23601	NP_037384	NM_013252	4D	219890_at
130	MGC3130	78995	NP_076937	NM_024032	4D	219879 s_at
42	FLJ13842	79698	NP_078921	NM_024645	4D	219877_at
2	ATF71P2	80063	NP_079273	NM_024997	4D	219870_at
	NARF	26502	NP_036468	NM_012336	4D	219862_s_at
	CLEC4E	26253	NP_055173	NM_014358	4D	219859_at
	ZNF14	7561	NP_066358	NM_021030	4D	219854_at
)	FLJ23506	79891	NP_079109	NM_024833	4D	219849_at
				NM_025174	4D	219846_at
	MTRFI	9617	NP_004285	NM_004294	4D	219822_at
	MRPS28	28957	NP 054737	NM_014018	4D	219819_s_at
	GPATC1	55094	NP_060495	NM_018025	4D	219818_s_at
	FLJ39616	51275	NP_057618	NM_016534	4D	219817_at
	VCIP135	80124	NP_079330	NM_025054	4D	219810_at
	FN5	56935	NP_064564	NM_020179	4D	219806 s at
	PILRA	29992	NP_038467	NM_013439	4D	219788_at
	ECT2	1894	NP_060568	NM_018098	4D	219787_s_at
	FLJ10996	54520	NP_061917	NM_019044	4D	219774_at
	GTDCI	79712	NP_078935	NM_024659	4D	219770_at
	B7H4	79679	NP_078902	NM_024626	4D	219768_at
	CRYZLI	9946	NP_005102	NM_005111	4D	219767_s_at
~	MGC4093	80776	NP_085055	NM_030578	4D	219766_at
6	FLJ12586	79673	NP_078896	NM_024620	4D	219765_at
	RPL36	25873	NP_056229	NM_015414	4D	219762 s at
8	FLJ21148	79918	NP_079136	NM_024860	4D	219751_at
	ZNF552	79818	NP_079038	NM_024762	4D	219741 x at
	PCDH9	5101	NP_065136	NM_020403	4D	219737_s at
İ	SIDTI	54847	NP_060169	NM_017699	4D	219734_at
				NM_014796	4D	219724 s_at

hypothetical protein El 199671	FLJ22671	79919	NP 079137	NM 024861	ŧ	220149 at
aldehyde dehydrogenase 8 family, member A1	ALDH8A1	64577	NP_072090	NM_022568	4D	220148 at
sorting nexin 11	SNXII	29916	NP_037455	NM_013323	4D	220140 s at
heart and neural crest derivatives expressed 1	HANDI	9421	NP_004812	NM_004821	4D	220138_at
solute carrier family 35, member F5	SLC35F5	80255	NP_079457	NM_025181	4D	220123_at
stabilin 2	STAB2	55576	NP_060034	NM_017564	4D	220114 s at
chromosome 9 open reading frame 39	C9orf39	54875	NP_060208	NM_017738	4D	220095_at
solute carrier family 2 (facilitated glucose trans	SLC2A6	11182	NP_060055	NM_017585	4D	220091_at
complement component 5 receptor 1 (C5a ligand)	C5R1	728	NP_001727	NM_001736	4D	220088_at
				NM_018391	4D	220079 s at
				NM_018391	4D	220078_at
preB lymphocyte gene 3	VPREB3	29802	NP_037510	NM_013378	4D	220068_at
melanoma antigen family C, 2	MAGEC2	51438	NP_057333	NM_016249	4D	220062_s_at
BCR downstream signaling 1	BRDG1	26228	NP_036240	NM_012108	4D	220059_at
interleukin 23, alpha subunit p19	IL23A	51561	NP_057668	NM_016584	4D	220054_at
TERF1 (TRF1)interacting nuclear factor 2	TINF2	26277	NP_036593	NM_012461	4D	220052 s at
sirtuin (silent mating type information regulation	SIRT4	23409	NP_036372	NM_012240	4D	220047_at
serum/glucocorticoid regulated kinaselike	SGKL	23678	NP_037389	NM_013257	4Đ	220038_at
lipocalininteracting membrane receptor	LIMR	55716	NP_060583	NM_018113	4D	220036_s_at
nucleoporin 210kDa	NUP210	23225	NP_079199	NM_024923	4D	220035 at
Ras interacting protein 1	RASIPI	54922	NP_060275	NM_017805	4D	220027 s_at
EROHike beta (S. cerevisiae)	EROILB	56605	NP_063944	NM_019891	4D	220012_at
peptidyl arginine deiminase, type IV	PADI4	23569	NP_036519	NM_012387	4D	220001 at
sialic acid binding Iglike lectin 5	SIGLECS	8778	NP_003821	NM_003830	4D	220000_at
				NM_018621	4D	219999_at
HSPC159 protein	HSPC159	29094	NP_054900	NM_014181	4D	219998_at
COP9 constitutive photomorphogenic homolog subunit	COPS7B	64708	NP_073567	NM_022730	4D	219997_s_at
interleukin 21 receptor	IL21R	50615	NP_068570	NM 021798	4D	219971 at
angiotensin I converting enzyme (peptidyldipeptida	ACE2	59272	NP_068576	NM_021804	4D	219962_at
UDPNacetylalphaDgalactosamine:polypeptide Nacetylg	GALNT6	11226	NP_009141	NM_007210	4D	219956_at
chromosome 11 open reading frame 17	CHOIL!	20072	141 000000	14141 OZOOZO	70	21/// 3 at

220418_at 220426_at 220419 s at 220376_at 220329 s_at 220282 at 20443 s at 220435_at 220404 at 220397_at 220391_at 220367 s at 220355 s at 220345 at 220341_s_at 220330_s_at 220327_at 220316 at 220283 at 220242 x_at 220241_at 220195 at 220176_at 220175 s_ar 220156 220261_s_at 220240 s at 20235 s at 220213_at 20212 s at 20206 at at 40 4D 40 4 4 4 4D 4 40 4 40 4 4 4 4 40 4 4 4D 40 40 4D 4D NM_012476 NM_018713 NM_024545 NM 018165 NM_024059 NM_013396 NM_01896 NM_014076 NM 020128 NM 024784 NM 022901 NM_024993 NM_016175 NM_022136 NM_017909 NM_018260 NM_017905 NM 017905 NM_016206 NM_022123 NM_024746 NM_024557 NM_018106 NM_018372 NM 018692 NM_022065 NM_024772 NM_018328 NM_025152 NM 020667 NM_024593 NP_076964 NP_036608 NP 061183 NP_037528 NP_061834 NP_064513 NP_079060 NP_075052 NP_078821 NP_060635 NP_079269 NP_057259 NP_071419 NP_060379 NP_057290 NP_071406 NP_079022 NP_078833 NP_060576 NP_060730 NP_060375 NP 060375 NP_060842 NP_071348 NP 079048 NP 060798 NP 079428 NP 078869 55532 64922 79025 53347 56890 55193 80059 64067 55146 29761 79842 79595 51149 64092 55005 79802 79608 55762 55002 55002 5579 63892 79645 389136 55777 80224 79830 USP25 VAX2 SLC30A10 MGC5356 PBI MDM SAP130 LRRTM4 LOC51149 RIFI UBASH3A ZBTB3 LRRC19 SAMSNI NPAS3 ZMYMI C6orf96 FLJ38507 FLJ13840 ZDHHC4 FLJ10891 C13orf11 Cl3orf1 THADA MBD5 NUBPL FLJ11767 ventral anterior homeobox 2 solute carrier family 30 (zinc transporter), membe hypothetical protein MGC5356 ubiquitin specific protease 25 ubiquitin associated and SH3 domain containing. Mdm4, transformed 3T3 cell double minute 1, p53 zinc finger and BTB domain containing 3 mSin3 Aassociated protein 130 leucine rich repeat transmembrane neuronal 4 truncated calcium binding protein leucine rich repeat containing 19 polybromo I RIC3 protein chromosome 13 open reading frame 11 chromosome 13 open reading frame 11 receptorinteracting factor I localisation si SAM domain, SH3 domain and nuclear chromosome 6 open reading frame 96 colon carcinoma related protein neuronal PAS domain protein 3 hypothetical protein FLJ13840 zinc finger, DHHC domain containing hypothetical protein FL11089 thyroid adenoma associated zinc finger, MYM domain containing 1 methylCpG binding domain protein 5 nucleotide binding proteinlike hypothetical protein FLJ1176

APPENDIX D

NY1-4308244v1

solute carrier family 9 (sodium/hydrogen	SLC9A5	6553	NP_004585	NM_004594	4D	220650_s_at
leucine rich repeat containing 31	LRRC31	79782	NP_079003	NM_024727	4D	220622_at
forkhead box E3	FOXE3	2301	NP 036318	NM_012186	40	220621 at
chromosome I open reading frame 42	Clorf42	54544	NP_061933	NM_019060	4D	220620_at
zinc finger protein 532	ZNF532	55205	NP_060651	NM_018181	4D	220617 s at
male sterility domain containing 1	MLSTDI	55711	NP_060569	NM_018099	4D	220615_s_at
leucine rich repeat (in FLII) interacting protein	LRRFIP2	9209	NP_006300	NM_006309	4D	220610 s at
ADPribosylationlike factor 6 interacting protein 4	ARL6IP4	51329	NP_061164	NM_018694	4D	220597 s at
uncharacterized hematopoietic stem/progenitor cell	MDS028	55846	NP_060933	NM_018463	4D	220590_at
				NM_025086	4D	220583_at
				NM_025071	4D	220582_at
resistin	RETN	56729	NP_065148	NM_020415	4D	220570_at
phosphoinositide3kinase, regulatory subunit 5, p10	PIK3R5	23533	NP_055123	NM_014308	4D	220566_at
panhematopoietic expression	PHEMX	10077	NP_005696	NM_005705	4D	220558_x_at
family with sequence similarity 35, member A	FAM35A	54537	NP_061927	NM_019054	4D	220547_s_at
chromosome 10 open reading frame 92	C10orf92	54777	NP_060079	NM_017609	4D	220539_at
myotubularin related protein 8	MTMR8	55613	NP_060147	NM_017677	4D	220537_at
				NM_024853	4D	220533_at
mitochondrial ribosomal protein L20	MRPL20	55052	NP_060441	NM_017971	4D	220526_s_at
deleted in liver cancer l	DLC1	10395	NP_079043	NM_024767	4D	220512_at
Tcomplex protein 1	CESK1	150160	NP_055221	NM_014406	4D	220508_at
hepcidin antimicrobial peptide	HAMP	57817	NP_066998	NM_021175	4D	220491_at
				NM_018565	4D	220490_at
				NM_017698	4D	220486_x_at
deafness locus associated putative guanine nucleot	DELGEF	26297	NP_036271	NM_012139	4D	220482_s_at
G proteincoupled receptor 75	GPR75	10936	NP_006785	NM_006794	4D	220481_at
hypothetical protein LOC55924	LOC55924	55924	NP_061972	NM_019099	4D	220476_s_at
myc target l	MYCT1	80177	NP_079383	NM_025107	4D	220471_s_at
				NM_025032	4D	220467_at
solute carrier organic anion transporter family, m	SLCOICI	53919	NP_059131	NM_017435	4D	220460_at
histamine receptor H3	HRH3	11255	NP_009163	NM_007232	4D	220447_at
carbohydrate (Nacetylglucosamine 6O) sulfotransfer	CHST4	10164	NP_005760	NM_005769	4D	220446_s_at

ring finger protein 170	RNF170	81790	NP_112216	NM_030954	4D	220985 s at
sprouty homolog 4 (Drosophila)	SPRY4	81848	NP_112226	NM_030964	4D	220983_s_at
erythrocyte membrane protein band 4.1 like 5	EPB41L5	57669	NP_065960	NM_020909	4D	220977 x at
keratin associated protein 11	KRTAP11	81851	NP_112229	NM_030967	4D	220976_s_at
interleukin 17E	IL17E	64806	NP_073626	NM_022789	4D	220971_at
				NM_030892	4D	220969 s at
ribosomal protein L22	RPL22	6146	NP_000974	NM_000983	4D	220960_x_at
MANSC domain containing 1	MANSCI	54682	NP_060520	NM_018050	4D	220945_x_at
				NM_025190	4D	220940_at
				NM_024084	4D	220934 s at
heparanase 2	HPSE2	60495	NP_068600	NM_021828	4D	220927 s at
G proteincoupled receptor 157	GPR157	80045	NP_079256	NM_024980	4D	220901_at
PR domain containing 12	PRDM12	59335	NP_067632	NM_021619	4D	220894 x at
DEAD (AspGluAlaAsp) box polypeptide 47	DDX47	51202	NP_057439	NM_016355	4D	220890_s_at
				NM_018614	40	220883 at
				NM_018575	4D	220874_at
				NM_014123	4D	220854 at
FERM domain containing 1	FRMDI	79981	NP_079195	NM_024919	4D	220819_at
gremlin 2 homolog, cysteine knot superfamily (Xeno	GREM2	64388	NP_071914	NM_022469	4D	220794_at
urotensin 2	UTS2	10911	NP_068835	NM_021995	4D	220785 at
urotensin 2	UTS2	10911	NP_068835	NM_021995	40	220784 s_at
potassium inwardlyrectifying channel, subfamily J,	KCNJ14	3770	NP_037480	NM_013348	4D	220776_at
dymeclin	DYM	54808	NP_060123	NM_017653	4D	220774_at
chromosome 6 open reading frame 48	C6orf48	50854	NP_058643	NM_016947	4D	220755_s_at
zinc finger protein 580	ZNF580	51157	NP_057286	NM_016202	4D	220748 s at
WD repeat domain 10	WDR10	55764	NP_060732	NM_018262	4D	220744 s at
inorganic pyrophosphatase 2	PPA2	27068	NP_008834	NM_006903	4D	220741_s_at
				NM_024992	4D	220715_at
				NM_018563	4D	220704_at
				NM_018616	4D	220702_at
KIAA1622	KIAA1622	57718	NP_066009	NM_020958	4D	220673 s at
exchanger						

calcium channel voltagedenandant germen	CACNGS	27091	NP 055219	NM 014404	45	221401_at
ectodysplasin A2 receptor	EDA2R	60401	NP_068555	NM_021783	4D	221399 at
				NM_020527	40	221379 at
recombining binding protein suppressor of hairless	RBPSUHL	11317	NP_055091	NM_014276	4D	221377 s at
motilin receptor	MLNR	2862	NP 001498	NM_001507	4D	221365_at
hypothetical protein dJ122O8.2	DJ12208.2	57226	NP_065199	NM_020466	4D	221311 x at
fibroblast growth factor receptor substrate 2	FRS2	10818	NP_006645	NM_006654	4D	221308_at
G proteincoupled receptor 22	GPR22	2845	NP_005286	NM_005295	4D	221288 at
pseudouridylate synthase 3	PUS3	83480	NP_112597	NM_031307	4D	221277 s at
				NM_030825	4D	221235_s_at
hypothetical protein FLJ20628	FLJ20628	55006	NP_060380	NM_017910	4D	221229_s_at
hypothetical protein FLJ23342	FLJ23342	79684	NP_078907	NM_024631	4D	221208 s at
YEATS domain containing 2	YEATS2	55689	NP 060493	NM_018023	4D	221203_s_at
chromosome 18 open reading frame 8	C18orf8	29919	NP_037458	NM_013326	4D	221190_s_at
				NM_025050	4D	221179_at
				NM_025043	4D	221177_at
histamine receptor H4	HRH4	59340	NP_067637	NM_021624	4D	221169_s_at
carbohydrate (Nacetylglucosamine 6O) sulfotransfer	CHST5	23563	NP_036258	NM_012126	ŧ	221104_X_at
achaetescute complex (Drosophila) homologlike 3	ASCL3	56676	NP_065697	NM_020646	4	221161 at
cell cycle progression 1	CCPG1	9236	NP_004739	NM_004748	40	221156_x_at
HT001 protein	HT001	28990	NP_054784	NM_014065	40	221135 s at
angiopoietin 4	ANGPT4	51378	NP_057069	NM_015985	4D	221134_at
hypothetical protein FLJ10826	FLJ10826	55239	NP_060703	NM_018233	4D	221090_s_at
family with sequence similarity 31, member C	FAM31C	79958	NP_079174	NM_024898	4D	221080_s_at
armadillo repeat containing 4	ARMC4	55130	NP_060546	NM_018076	4D	221077_at
spermatogenesis associated 1	SPATAI	64173	NP 071749	NM_022354	4D	221057_at
anterior pharynx defective 1 homolog B (C. elegans	APHIB	83464	NP_112591	NM_031301	4D	221036_s_at
transmembrane protease, serine 5 (spinesin)	TMPRSS5	80975	NP_110397	NM_030770	4D	221032 s at
RAB33B, member RAS oncogene family	RAB33B	83452	NP_112586	NM_031296	4D	221014 s_at
chromosome I open reading frame 14	Clorf14	81626	NP_112195	NM_030933	4D	220996 s at
likely ortholog of rat vacuole membrane protein 1	VMPI	81671	NP_112200	NM_030938	4D	220990 s_at
tigger transposable element derived 6	TIGD6	81789	NP_112215	NW USOSS	ť	220700 S at

likely ortholog of mouse synembryn	RIC8	60626	NP_068/51	NM 021932	=	5 (10H) 5 at
arginineglutamic acid dipeptide (RE) repeats	RERE					2 2
syntaxin 16	STX16	5	433	433		
uncharacterized hypothalamus protein HT007	HT007		NP 060950			
signaltransducing adaptor protein2	STAP2	55620	NP_060190			
HSPC171 protein	HSPC171	29100	NP_054906			
histone 3, H2a	HIST3H2A	92815	NP 254280			
sterol Oacyltransferase (acylCoenzyme A: cholester	SOATI	0040	NP_005092			É
lymphoid enhancerbinding factor 1	LEFI	51176	NP_057353			221561 or
contain	CIVIOI LDZ	0.10	1.000			
general transcription factor IIH, polypeptide 2, 4	CBISBI DO	2716	NP 113664	NM 031476		221541 at
nypothetical protein FLJ11301	L F J L 1901	14000	NID 001500			221540 v at
ankyrın repeat domain 27 (VPS9 domain)	ANKKU2/	640/9	NP 060855	NM 018385	4D 6	221536 s at
cell division cycle associated 8	CDCA8	55143	NP 0605/1			221320 s at
cofactor required for Sp1 transcriptional activati	CRSP6	9440	NP_004259			221520 - at
hypothetical protein FLJ20232	FLJ20232	54471	NP_061881			221515 s at
cell cycle progression 1	CCPG1	9236	NP_065790	NM_020739		221511 x at
karyopherin alpha 3 (importin alpha 4)	KPNA3	3839	NP_002258	NM_002267		221503_s_at
eukaryotic translation initiation factor 3, subuni	EIF3S12	27335	NP_037366	NM_013234		221494_x_at
sprouty homolog 4 (Drosophila)	SPRY4	81848	NP_112226			221489 S at
galactosyltransferase,	DIGNET	1001				
galaciosyliransierase,	DACALTZ	033/	ND 004767	NM 004776	4	221485 at
UDPGal:betaGlcNAc beta 1,4	B4GALT5	9334	NP_004767	NM_004//6	‡	721404 at
BCL2/adenovirus E1B 19kDa interacting protein 31ik	BNIP3L	000	NF_004322	MM 004331		200
glucose6phosphatase, catalytic, 2	G6PC2	57818	NP 066999	NM 021170		221475 at
olfactory receptor, family 2, subfamily W, member	OK2W1	26007	M112100	NIM GOODS		221452 01
Tcell immunomodulatory protein	CDA08	81533	NP 110417	NM 030003	40	221451 s at
chromosome 14 open reading frame 156	C14orf156	81892	NP 112487	NM 031210		221424 S at
cyclin L2	CCNL2	81669	NP 112199	NM 030937		221424 S at
subunit 5						77177

contour commolile feetle	CTGLF1	119016	NP 597703	NM 133446	4D	2219/1 x at
homolog, yea						
ubiquitinconjugating enzyme E2H (UBC8	UBE2H	7328	NP_003335	NM_003344	4D	221962_s_at
hypothetical protein LOC283232	LOC283232	283232	NP_777600	NM_174940	4D	221951_at
chromosome 14 open reading frame 87	C14orf87	51218	NP_057501	NM_016417	4D	221932 s_at
hypothetical protein DKFZp76112123	DKFZp76112123	83637	NP_777589	NM_174929	4D	221924_at
hypothetical protein MGC1203	MGC1203	79140	NP_077272	NM_024296	4D	221912 s_at
likely ortholog of mouse hypoxia induced gene 1	HIGI	25994	NP_054775	NM_014056	4D	221896 s_at
ecotropic viral integration site 1	EVII	2122	NP_005232	NM_005241	40	221884 at
major histocompatibility complex, class I, F	HLAF	3134	NP_061823	NM_018950	4D	221875 x at
TBC1 domain family, member 12	TBC1D12	23232	XP_051081	XM_051081	4D	221858 at
plakophilin I (ectodermal dysplasia/skin fragility	PKP1	5317	NP_001005337	NM_001005337	4D	221854 at
centaurin, gammalike family, member 1	CTGLF1	119016	NP_597703	NM_133446	4D	221850_x_at
RAP2A, member of RAS oncogene family	RAP2A	5911	NP_066361	NM_021033	4D	221830_at
hypothetical protein FLJ20436	FLJ20436	54934	NP_060292	NM_017822	4D	221821_s_at
KIAA1698 protein	KIAA1698	80789	NP_085131	NM_030628	4D	221818_at
PHD finger protein 11	PHF11	51131	NP_057203	NM_016119	4D	221816 s at
RAB9A, member RAS oncogene family	RAB9A	9367	NP_004242	NM_004251	4D	221808_at
nuclear receptor binding factor 2	NRBF2	29982	NP_110386	NM_030759	4D	221803_s_at
ribosomal protein L22	RPL22	6146	NP_000974	NM_000983	4D	221775_x_at
jumonji domain containing 1C	JMJD1C	221037	NP_004232	NM_004241	4D	221763_at
YTH domain family, member 1	YTHDFI	54915	NP_060268	NM_017798	4D	221741_s_at
collagen, type V, alpha 2	COL5A2	1290	NP_000384	NM_000393	4D	221729_at
Ctype lectin domain family 4, member A	CLEC4A	50856	NP_057268	NM_016184	4D	221724 s at
solute carrier family 4, sodium bicarbonate cotran	SLC4A5	57835	NP_067019	NM_021196	Đ	221723_s_at
hypothetical protein FLJ10439	FLJ10439	54663	NP_060563	NM_018093	4D	221712 s at
hypothetical protein FLJ12750	FLJ12750	79720	NP_078943	NM_024667	4D	221704_s_at
BBPlike protein 2	BLP2	80213	NP_510883	NM_078474	4D	221702_s_at
ubiquitin A52 residue ribosomal protein fusion pro	UBA52	7311	NP_003324	NM_003333	4D	221700_s_at
mitochondrial ribosomal protein L34	MRPL34	64981	NP_076426	NM_023937	1 D	221692 s at
Down syndrome critical region gene 5	DSCR5	51227	NP_710149	NM_153682	4D	221689 s at
abhydrolase domain containing 6	ABHD6	57406	NP_065727	NM_020676	4D	221679 s at
coronin, actin binding protein, 1C	CORO1C	23603	NP_055140	NM_014325	4D	221676 s at
LIM nomeobox 3	LIIAS	0022	141 000017			

feetic expressed comence 27	TEX27	60685	NP_068762	NM_021943	4D	222493 s_at
ubiquitinconjugating enzyme E2Q (putative)	UBE2Q	55585	NP_060052	NM_017582	45	222480 at
translocase of outer mitochondrial membrane 22 hom	TOMM22	36993	NP_064628	NM_020245	ŧ	2224)+ 5 at
mediator of RNA polymerase II transcription, subun	MED4		NP_054885	NW 014160	ŧ	22.477. at
vacuolar protein sorting 24 (yeast)	VPS24		NP 057163	NM 0160/9	ŧ	22430 S at
coronin, actin binding protein, 1C	CORO1C	23603	NP_055140	NM_014325	5	222409 at
yippeelike 5 (Drosophila)	YPEL5	51646	NP_057145	NM_016061	40	222408 s_at
butyrateinduced transcript 1	HSPC121	51495	NP_057479	NM_016395	4D	222404 x at
mitochondrial carrier homolog 2 (C, elegans)	MTCH2	23788	NP 055157	NM_014342	4D	222403 at
small membrane protein i	SMPI	23585	NP_055128	NM_014313	4D	222401 s at
hematological and neurological expressed	INH	51155	NP_057269	NM_016185	4D	222396_at
programmed cell death 6 interacting protein	PDCD6IP	10015	NP_037506	NM_013374	4D	222394 at
CG130 protein	CG130	51611	NP_057042	NM_015958	4D	222360_at
splicing factor, arginine/serinerich 15	SFRS15	57466	NP_065757	NM_020706	4D	222311_s_at
sirtuin (silent mating type information regulation	SIRT4	23409	NP_036372	NM_012240	4	222248 s at
paired immunoglobinlike type 2 receptor alpha	PILRA	29992	NP_840057	NM_1/82/3	ŧ	722218 S at
LAG1 longevity assurance homolog 2 (S. cerevisiae)	LASS2	29956	NP_037516	NM_013384	ŧ	222216
hypothetical protein FLJ22104	FLJ22104	65084	NP_075069	NM 022918	ŧ	222212 at
cell cycle progression 1	CCPG1	9236	NP_065790	NM_020739	ŧ	222156 x at
G proteincoupled receptor 89	GPR89	51463	NP_057418	NM_016334	4D	222140 s at
Daspartate oxidase	DDO	8528	NP_004023	NM_004032	4D	222134_at
FtsJ homolog 2 (E. coli)	FTSJ2	29960	NP_037525	NM_013393	4D	222130 s at
amphoterin induced gene 2	AMIGO2	347902	NP_862830	NM_181847	4D	222108 at
activating transcription factor	ATF1	466	NP_005162	NM_005171	4D	222103_at
solute carrier family 2 (facilitated glucose trans	SLC2A3	6515	NP_008862	NM_006931	4D	222088 s_at
interleukin 27 receptor, alpha	IL27RA	9466	NP_004834	NM_004843	4D	222062_at
hypothetical protein MGC3731	MGC3731	79159	NP 077289	NM_024313	4D	222057 at
zinc finger protein 45	ZNF45	7596	NP_003416	NM_003425	4D	222028_at
leucinerich repeats and calponin homology (CH) dom	LRCH4	4034	NP_002310	NM_002319	ŧ	22201/_X_at
tcomplex 1	TCP1	6950	NP_110379	NM_030752	ŧ	222017 at
hypothetical protein FLJ10534	FLJ10534	55/20	06C000 JN	[A A 010120	1	222010

HSPC128 protein	HSPC128	29080	NP_054886	NM_014167	4D	222792 s at
signal peptidase complex subunit 3 homolog (S. cer	SPCS3	60559	NP_068747	NM_021928	4D	222753_s_at
sex comb on midleglike 1 (Drosophila)	SCML1	6322	NP_006737	NM_006746	4D	222747 s at
chromosome 15 open reading frame 29	C15orf29	79768	NP_078989	NM_024713	4D	222745_s_at
trimethyllysine hydroxylase, epsilon	TMLHE	55217	NP_060666	NM_018196	4D	222744 s at
hypothetical protein FLJ23375	FLJ23375	80021	NP_079232	NM_024956	4D	222739_at
hypothetical protein MGC5306	MGC5306	79101	NP_077021	NM_024116	4D	222728_s_at
lactamase, beta 2	LACTB2	51110	NP_057111	NM_016027	4D	222714 s at
postsynaptic protein CRIPT	CRIPT	9419	NP_054890	NM_014171	4D	222702 x at
x 010 protein	MDS010	56983	NP_689518	NM_152305	4D	222681_at
RP42 homolog	RP42	54165	NP_065691	NM_020640	4D	222678_s_at
BAllassociated protein 2like 1	BAIAP2L1	55971	NP_061330	NM_018842	4D	222676_at
vmar musculoaponeurotic librosarcoma oncogene homo	MALD	9933	W_003432	NW 003461	ŧ	222010_S_at
hypothetical protein FLJ11011	FLJ11011	55284	NP_001001481	NM_001001481	#	222657_s_at
hypothetical protein FLJ11011	FLJ11011	55284	NP_001001481	NM_001001481	4D	222656_at
trichorhinophalangeal syndrome I	TRPS1	7227	NP_054831	NM_014112	4D	222651 s at
exportin 4	XPO4	64328	NP_071904	NM_022459	4D	222649_at
hypothetical protein FLJ22662	FLJ22662	79887	NP_079105	NM_024829	4D	222639 s_at
mediator of RNA polymerase II transcription, subun	MED28	80306	NP_079481	NM_025205	4D	222636_at
phosphatidylinositol 4kinase type11 beta	PI4K2B	55300	NP_060793	NM_018323	4D	222631_at
nudE nuclear distribution gene E homolog 1 (A. nid	NDEI	54820	NP_060138	NM_017668	4D	222625_s_at
chromosome 10 open reading frame 84	C10orf84	63877	NP_071346	NM_022063	4D	222617_s_at
exosome component 1	EXOSC1	51013	NP_057130	NM_016046	4D	222609 s_at
xenotropic and polytropic retrovirus receptor	XPR1	9213	NP_004727	NM_004736	4D	222581_at
mitogenactivated protein kinase kinase kinase kina	MAP4K4	9448	NP_004825	NM_004834	4D	222547_at
zinc finger protein 395	ZNF395	55893	NP_061130	NM_018660	4D	222536 s at
chromosome 14 open reading frame 108	C14orf108	55745	NP_060699	NM_018229	4D	222531_s_at
ankyrin repeat and SOCS boxcontaining 6	ASB6	140459	NP_060343	NM_017873	4D	222524 s at
adaptorrelated protein complex 3, mu 1 subunit	AP3M1	26985	NP 996895	NM_207012	4D	222516 at
NEDD8 ultimate buster!	NYRENI8	51667	NP_057202	NM_016118	4D	222512_at
chromosome 7 open reading frame 2	C7orf2	64327	NP_071903	NM_022458	4D	222505_at

223017_at	22301	22300	22300	22300	22299	22299	222990_at	22298	222985_at	222984_at	22298		222969_at	2229	222949_at	2229:	22289	22289	22289	222875	22280	2228:	2228-	222830_at	222817_at	222805_at	222798 at	22279
7_at	223011_s_at	223008_s_at	223006 s_at	223004_s_at	222997_s_at	222992_s_at	0 at	222989 s at	S_at	84 at	222983 s at		9 at	222955 s_at	19 at	222934_s_at	222895_s_at	222893_s_at	222891_s_at	75_at	222867_s_at	222858_s_at	222847 s at	0 at	7_at)5 at)8 at	222794_x_at
4D	4D	4D	4D	4D	4D	4D	4D	4D	45	4D	4D		₽ D	4D	4D	4D	4D	4D	4D	4D	4D	4D	4D	4D	4D	4D	4D	4D
NM_015913	NM_017830	NM_032012	NM_032012	NM_016589	NM_018997	NM_005005	NM_013438	NM_013438	NM_012479	NM_016480	NM_016480		NM_020981	NM_207009	NM_022052	NM_014358	NM_022898	NM_024813	NM_022893	NM_020162	NM_016060	NM_014395	NM_022073	NM_198182	NM_025193	NM_024641	NM_001001484	NM_018109
NP_056997	NP_060300	NP_114401	NP_114401	NP_057673	NP_061870	NP_004996	NP_038466	NP_038466	NP_036611		NP_057564		NP_066191	NP_996892	NP_071335	NP_055173	NP_075049	NP_079089	NP_075044	NP_064547	NP_057144	NP_055210	NP_071356	NP_937825	NP_079469	NP 078917	NP_001001484	NP_060579
51060	54940	23731	23731	51300	54460	4715	29979	29979	7532	51247	51247		8708	404636	56000	26253	64919	79871	53335	56919	51003	27071	112399	29841	80270	79694	9317	55149
TLP19	OCIADI	C9orf5	C9orf5	C3orf1	MRPS21	NDUFB9	UBQLNI	UBQLNI	YWHAG	PAIP2	PAIP2		B3GALT1	FAM45A	NXF3	CLEC4E	BCLIIB	FLJ13150	BCLIIA	DHX33	MED31	DAPPI	EGLN3	TFCP2L2	HSD3B7	MANEA	PTER	PAPDI
endoplasmic reticulum thioredoxin superfamily memb	OCIA domain containing 1	chromosome 9 open reading frame 5	chromosome 9 open reading frame 5	chromosome 3 open reading frame 1	mitochondrial ribosomal protein S21	NADH dehydrogenase (ubiquinone) I beta subcomplex,	ubiquilin 1	ubiquilin l	tyrosine 3monooxygenase/tryptophan 5monooxygenase	poly(A) binding protein interacting protein 2	poly(A) binding protein interacting protein 2	1,3galactosyltransferase, p	UDPGal:betaGlcNAc beta	family with sequence similarity 45, member A	nuclear RNA export factor 3	Ctype lectin domain family 4, member E	Bcell CLL/lymphoma 11B (zinc finger protein)	hypothetical protein FLJ13150	Bcell CLL/lymphoma 11A (zinc finger protein)	DEAH (AspGluAlaHis) box polypeptide 33	mediator of RNA polymerase II transcription, subun	dual adaptor of phosphotyrosine and 3phosphoinosit	egl nine homolog 3 (C. elegans)	transcription factor CP2like 2	hydroxydelfa5steroid dehydrogenase, 3 beta and ste	mannosidase, endoalpha	phosphotriesterase related	PAP associated domain containing 1

cingulin	CGN	57530	NP_065821	NM 020770	4D	223233_s_at
ARV1 homolog (yeast)	ARV1	64801	NP_073623	NM_022786	4D	223223_at
zinc fingers and homeoboxes 1	ZHX1	11244	NP_009153	NM_007222	4D	223214 s_at
zinc finger, DHHC domain containing 16	ZDHHC16	84287	NP_932163	NM_198046	4D	223212_at
SWI/SNFrelated, matrixassociated actindependent re	SMARCADI	56916	NP_064544	NM_020159	4D	223197_s_at
sestrin 2	SESN2	83667	NP_113647	NM_031459	4D	223195_s_at
chromosome 6 open reading frame 69	C6orf69	222658	NP_775833	NM_173562	4D	223176_at
ubiquitin specific protease 25	USP25	29761	NP_037528	NM_013396	4D	223167_s_at
zinc finger, C3HC type 1	ZC3HC1	51530	NP_057562	NM_016478	4D	223163 s_at
chromosome 4 open reading frame 14	C4orf14	84273	NP_115689	NM_032313	4D	223157_at
hypothetical protein MGC2714	MGC2714	84259	NP_115675	NM_032299	4D	223151_at
DEAH (AspGluAlaHis) box polypeptide 36	DHX36	170506	NP_065916	NM_020865	4D	223139 s at
androgeninduced 1	AIG1	51390	NP_057192	NM_016108	4D	223136_at
hypothetical protein H17	H17	55572	NP_060017	NM_017547	4D	223128_at
secreted frizzledrelated protein 2	SFRP2	6423	NP_003004	NM_003013	4D	223121_s_at
hypothetical protein HSPC196	HSPC196	51524	NP_057548	NM_016464	4D	223113_at
ADPribosylhydrolase like 2	ADPRHL2	54936	NP_060295	NM_017825	4D	223097_at
mitochondrial ribosomal protein L51	MRPL51	51258	NP_057581	NM_016497	4D	223086_x_at
ring finger protein 19	RNF19	25897	NP_056250	NM_015435	4D	223085_at
cyclin Dtype bindingprotein 1	CCNDBP1	23582	NP_036274	NM_012142	4D	223084 s_at
hypothetical protein MGC2941	MGC2941	79142	NP_077273	NM_024297	4D	223081_at
NOL1/NOP2/Sun domain family, member 2	NSUN2	54888	NP_060225	NM_017755	4D	223076_s_at
hypothetical protein FLJ14753	FLJ14753	84641	NP_115947	NM_032558	4D	223073_at
selenoprotein K	SELK	58515	NP_067060	NM_021237	4D	223070_at
STARD3 Nterminal like	STARD3NL	83930	NP_114405	NM_032016	4D	223065_s_at
hypothetical protein LOC51255	LOC51255	51255	NP_057578	NM_016494	4D	223064_at
exportin 5	XPOS	57510	NP_065801	NM_020750	4D	223057 s at
HSPC182 protein	HSPC182	29101	NP_054907	NM 014188	4D	223053 x at
solute carrier family 40 (ironregulated transporte	SLC40A1	30061	NP_055400	NM 014585	4D	223044 at
Nacetyltransferase 5 (ARD1 homolog, S. cerevisiae)	NAT5	51126	NP_852669	NM_181528	4D	223040_at
vacuolar protein sorting 29 (yeast)	VPS29	51699	NP_476528	NM_057180	4D	223026 s at
likely ortholog of mouse nin one binding protein	NOBIP	28987	NP_054781	NM_014062	4D	223018_at

NP 036312 26296 NP 061326 25967 NP 061326 25967 NP 061326 25632 NP 060239 94103 NP 04329 54758 25970 NP 115665 85027 NP 115658 85027 NP 1156479 84273 NP 115649 84273 NP 115649 84233 NP 079001 79780 84273 NP 06528 50481 NP 066996 65977 57616 NP 067907 57616 NP 067908 29950 NP 116225 85025 NP 116225 85025 NP 165236 85025 NP 165256 85025	cappuccino	CNO	55330	NP_060836	NM_018366	4D	223431_at
NP_036312 20269 FBXO8	hypothetical protein FLJ20425		55646	NP_060286	NM 017816	4D	223413 s at
NP_036312 20269 FBXO8	kelch repeat and BTB (POZ) domain containing 7		84078	NP 115514	NM_032138	4D	223412_at
NP_063512 26269 FBXO8	chromosome 7 open reading frame 35		85025	NP_116325	NM_032936	4D	223396_at
NP_036312 20269 FINOS NP_061326 25967 DAPI3 NP_061326 25967 DAPI3 NP_115662 84246 MGC3399 NP_060239 35652 KIAA1333 NP_064809 94103 ORMDL3 NP_065758 82027 NID67 NP_057581 57266 KLPI NP_075781 57265 MGCC4289 NP_115633 83607 MGC4288 NP_115633 83607 MGC4288 NP_075791 57281 NTSC3 NP_075791 79780 FL23318 NP_075791 P12331 DKF2686C192 NP_05866 10283 DKF2686C192 NP_060964 55869 HDAC8 NP_060965 53869 HDAC8 NP_061964 65977 PLEKHA3 NP_061964 65977 PLEKHA3 NP_063907 57616 ZNE537 NP_063907 57616 ZNE537 NP_063907 57616 ZNE537	SERTA domain containing 1	SERTADI	29950	NP_037508	NM_013376	4D	223394_at
NP_036312 20269 FBXO8	zinc finger protein 537	ZNF537	57616	NP_065907	NM_020856	40	223393 s at
NP 036312 S269 FBXO8	zinc finger protein 537	ZNF537	57616	NP_065907	NM_020856	4D	223392 s at
NP 036312 20269 FBXO8 NP 061326 25967 DAPI3 NP 115662 84246 MGC5309 NP 060239 55652 KIAA1333 NP 060511 57106 KLPI NP 065758 84275 NID67 NP 057584 51260 CXorf26 NP 115633 82607 MGC4288 NP 115649 84275 MGC4288 NP 115649 84233 DKF2686C192 NP 0657075 22931 RAB18 NP 0656528 50484 RRAD2B NP 060956 53899 HDAC8 NP 1661964 65977 PLEKHA3	tripartite motifcontaining 4	TRIM4	89122	NP_149082	NM_033091	4D	223384_s_at
NP_063312 26269 FBXO8 NP_061326 53967 DAPI	pleckstrin homology domain containing, family A	PLEKHA3	65977	NP_061964	NM_019091	4D	223370_at
NP_036312 26269 FBXO8 NP_061326 25967 DAPI3 NP_061326 25967 DAPI3 NP_115662 84246 MGC3309 NP_060239 25652 KIAA1333 NP_644809 94103 ORMIDL3 NP_054758 28970 PTD012 NP_166565 85027 NID677 NP_065111 57106 KLPI NP_057584 51260 CXort26 NP_115691 84275 MGC4289 NP_115691 84275 MGC4289 NP_115691 84275 MGC4288 NP_115691 84275 MGC4288 NP_115691 84275 MGC4288 NP_115691 84275 MGC4288 NP_157573 21251 NTSC3 NP_079001 79780 FL23518 NP_079001 79780 FL23518 MGC4288 NP_15767 MGC4288 NP_079001 79780 FL23518 NP_079001 79780 FL23518 NP_079001 NP_	melanoma associated antigen (mutated) I	MUMI	84939	NP_116242	NM_032853	4D	223347_at
NP_036312 26269 FBXO8 NP_061326 55967 DAPI 3 NP_061326 55967 DAPI 3 NP_115662 55632 KIAA1333 NP_060239 55632 KIAA1333 NP_054758 52870 NTD012 NP_05478 57106 KLPI TD012 NP_05478 57106 KLPI TD012 NP_057584 51260 CXort26 NP_157584 51260 CXort26 NP_157584 51260 CXort26 NP_157584 51260 KICPI 3 NP_057573 51251 MGC4268 NP_157584 51251 MGC4268 NP_157584 51251 MGC4268 NP_157584 51251 MGC4268 NP_057573 51251 MGC4268 NP_057573 51251 KABI8 NP_057075 22931 KABI8 NP_057075 22931 KABI8 NP_057075 23934 KRNEB	histone deacetylase 8	HDAC8	55869	NP_060956	NM_018486	4D	223345_at
NP 036312 26269 EBXO8	ribonucleotide reductase M2 B (TP53 inducible)	RRM2B	50484	NP_056528	NM_015713	4D	223342_at
NP_036312 26269 FBXO8 NP_061326 25967 DAPI3 NP_061326 25967 DAPI3 NP_115662 84246 MGC5399 NP_060239 25652 KIAA1333 NP_060239 25652 KIAA1333 NP_060239 25652 KIAA1333 NP_060239 25652 KIAA1333 NP_054758 28970 PITD012 NP_05656 88207 NID012 NP_05758 85077 KIAPI NP_05758 81250 CXort26 NP_115633 81267 MGC4399 NP_115633 81267 MGC4288 NP_07573 21251 NTSC3 NP_07579 179780 FL23518 NP_075901 179780 NP_075901 NP_075901 179780 NP_075901 179780 NP_075901 179780 NP	ATPase inhibitory factor 1	ATPIFI	93974	NP_835497	NM_178190	4D	223338 s at
NP_063312 26269 FBXO8 NP_061326 55967 DAPI 3 NP_061326 55967 DAPI 3 NP_1060239 55652 KIAA1333 NP_060239 55652 KIAA1333 NP_060239 55652 KIAA1333 NP_064738 58970 PTD012 NP_16668 58970 PTD012 NP_16568 58207 NID67 NP_057581 57106 KLPI KLPI 3 NP_057581 51260 KALPI 3 NP_115691 54275 MGC4268 NP_1057533 51251 MGC4268 NP_1057533 51251 MTSC3 NP_057533 51251 MTSC3 NP_057533 51251 MTSC3 NP_057533 51251 MTSC3 NP_057501 S1251 MTSC3 NP_057573 S1251 MTSC3 NP_057573 S1251 MTSC3 NP_057573 S1251 MTSC3 NP_057573 S1251 MTSC3 MTSC3 NP_057573 S1251 MTSC3 MTSC3 NP_057573 S1251 MTSC3 MTSC3 NP_057573 S1251 MTSC3 MTSC3 NP_057573 MTSC3	serologically defined colon cancer antigen 10	SDCCAG10	10283	NP_005860	NM_005869	4D	223337_at
NP_036312 26269 FBX08 NP_061326 55967 DAP13 NP_115662 84246 MGC3309 NP_060239 55652 KIAA1333 NP_060239 55652 KIAA1333 NP_044809 94103 ORMDL3 NP_054758 28970 PTD012 NP_051758 28970 PTD012 NP_06511 57106 KLP1 NP_05754 51260 CXort26 NP_115643 84275 MGC4399 NP_115643 84275 MGC4399 NP_057573 51251 NT5C3 NP_115649 84233 DKF25886C192	RAB18, member RAS oncogene family	RAB18	22931	NP_067075	NM_021252	4D	223336 s at
NP_063312 26269 FBXO8 NP_061326 55967 DAP13 NP_061326 55967 DAP13 NP_15662 58622 KIAA1333 NP_060239 55652 KIAA1333 NP_060239 55652 KIAA1333 NP_060239 55652 KIAA1333 NP_064738 94103 ORMDL3 NP_054738 94103 ORMDL3 NP_054738 85027 NID67 NP_16563 85027 NID67 NP_07584 51260 CXort26 NP_075734 51250 KIZP1 NP_075737 51251 NITSC3 NP_075737 51251 NITSC3 NP_075901 79780 FL232518	hypothetical protein DKFZp586C1924		84233	NP_115649	NM_032273	Đ	223334_at
NP_063312 26269 FBX088 NP_061326 25967 DAP13 NP_115662 84246 MGC3399 NP_060239 55662 K1AA1333 NP_060239 55662 K1AA1333 NP_060239 55662 K1AA1333 NP_06124809 94103 ORMDL3 NP_054788 28970 PTD012 NP_05111 57106 KLP1 NP_05784 51260 CXort26 NP_115693 83027 MGC4399 NP_115693 83067 MGC4399 NP_057573 51251 MTCG3	hypothetical protein FLJ23518	FLJ23518	79780	NP 079001	NM_024725	4D	223301 s at
NP_063312 26269 FBX088 NP_061326 25967 DAPI 3 NP_061326 25967 DAPI 3 NP_115662 84246 MGC3309 NP_060239 25652 KIAA1333 NP_060239 25652 KIAA1333 NP_064809 94103 ORMIDL3 NP_054738 28970 PTD012 NP_105656 88027 ND67 NP_057544 251260 CXort26 NP_115633 83607 MGC4399 NP_115633 83607 MGC4268	5'nucleotidase, cytosolic III	NT5C3	51251	NP_057573	NM_016489	4D	223298_s_at
NP_063312 26269 FBXO8 NP_061326 55967 DAP13 NP_061326 55967 MGCC3309 NP_060239 55632 KIAA1333 NP_060239 55632 KIAA1333 NP_060239 55632 KIAA1333 NP_0641788 28970 PTD012 NP_0641788 28970 PTD012 NP_065111 57106 KLP1 NP_057584 51260 CXord26 NP_057584 51260 CXord26	hypothetical protein MGC4268	MGC4268	83607	NP_113633	NM_031445	4D	223297_at
NP_063312	mitochondrial carrier protein	MGC4399	84275	NP_115691	NM_032315	4D	223296_at
NP_063112 26269 FBX08 NP_061326 55967 DAP13 NP_115662 84246 MGC5309 NP_060239 55652 KIAA1333 NP_060239 55652 KIAA1333 NP_064809 94103 ORMDL3 NP_054738 28970 PTD012 NP_06511 57106 KLP1	chromosome X open reading frame 26	CXorf26	51260	NP_057584	NM_016500	4D	223294_at
NP_063312 26269 FBXO8 NP_061326 55967 DAP13 NP_061326 55967 MGCC3309 NP_060239 55632 KIAA1333 NP_060239 55632 KIAA1333 NP_0641738 28970 PTD012 NP_164565 88027 NID67	K562 cellderived leucinezipperlike protein 1	KLPI	57106	NP_065111	NM_020378	4D	223284_at
NP_063312 26269 FBX08 NP_061326 53567 DAP13 NP_115662 84246 MGC5369 NP_060239 55632 KIAA1333 NP_060239 55632 KIAA1333 NP_644809 94103 ORMDI3 NP_644788 28970 PTD012	putative small membrane protein NID67	NID67	85027	NP_116565	NM_032947	4D	223276_at
NP_06312	PTD012 protein	PTD012	28970	NP_054758	NM_014039	4D	223268_at
NP_036312 26269 FBXO8 NP_061326 35967 DAP13 NP_1560239 35262 MGCC3309 NP_060239 353622 KIAA1333 NP_060239 353622 KIAA1333	ORMHike 3 (S. cerevisiae)	ORMDL3	94103	NP_644809	NM_139280	4D	223259_at
NP_06132 26269 FBXO8 NP_061326 535967 DAP13 NP_115602 84246 MGCS399 NP_060239 55632 KIAA1333	KIAA1333	KIAAI333	55632	NP_060239	NM_017769	4D	223256_at
NP_063312 26269 FBX08 NP_061326 55967 DAP13 NP_115662 84246 MGC53399	KIAA1333	KIAA1333	55632	NP_060239	NM_017769	4D	223255_at
NP_036312 26269 FBXO8 NP_061326 55967 DAP13	hypothetical protein MGC5309	MGC5309	84246	NP_115662	NM_032286	4D	223247_at
NP_036312 26269	13kDa differentiationassociated protein	DAP13	55967	NP_061326	NM_018838	4D	223244 s at
ı	Fbox protein 8	FBXO8	26269	NP_036312	NM_012180	4D	223240_at
NP 115517 84081	DKFZP434K142 hypothetical protein DKFZp434K1421	DKFZP434K142 1	84081	NP_115517	NM_032141	4D	223236_at

thymocyte protein thy28	THY28	29087	NP_054893	NM_014174	4D	223711_s_at
winglesstype MMTV integration site family, member	WNT10A	80326	NP_079492	NM_025216	4D	223709_s_at
chromosome 22 open reading frame 23	C22orf23	84645	NP_115950	NM_032561	4D	223706 at
latent transforming growth factor beta binding pro	LTBP2	4053	NP_000419	NM_000428	4D	223690_at
hypothetical protein MGC11102	MGC11102	84285	NP_115701	NM_032325	4D	223682 s at
CDC42 small effector 1	CDC42SE1	56882	NP_064624	NM_020239	4D	223674 s at
CG130 protein	CGI30	51611	NP_057042	NM_015958	4D	223671_x_at
hemogen	HEMGN	55363	NP_060907	NM_018437	4D	223670_s_at
nuclear receptor binding factor 2	NRBF2	29982	NP_110386	NM_030759	4D	223650_s_at
Jtype cochaperone HSC20	HSC20	150274	NP_741999	NM_172002	4D	223647 x at
zinc ribbon domain containing, 1	ZNRDI	30834	NP_055411	NM_014596	4D	223639 s_at
RASD family, member 2	RASD2	23551	NP_055125	NM_014310	4D	223634 at
family with sequence similarity 14, member A	FAM14A	83982	NP_114425	NM_032036	4D	223626_x_at
G proteincoupled receptor 34	GPR34	2857	NP_005291	NM_005300	4D	223620_at
ropporin 1like	ROPNIL	83853	NP_114122	NM_031916	4D	223609_at
kelch repeat and BTB (POZ) domain containing	KBTBD2	25948	NP_056298	NM_015483	4D	223584 s_at
hypothetical protein FLJ23467	FLJ23467	79626	NP_078851	NM_024575	4D	223583_at
chromosome 6 open reading frame 203	C6orf203	51250	NP_057571	NM_016487	4D	223576_at
chromosome I open reading frame 26	Clorf26	54823	NP_060143	NM_017673	4D	223548_at
ankyrin repeat domain 32	ANKRD32	84250	NP_115666	NM_032290	4D	223542_at
nudix (nucleoside diphosphate linked moiety X)type	NUDT12	83594	NP_113626	NM_031438	4D	223535_at
G proteincoupled receptor 89	GPR89	51463	NP_057418	NM_016334	4Đ	223531_x_at
FLJ20859 gene	FLJ20859	64745	NP_073571	NM_022734	4D	223528 s_at
chromosome 9 open reading frame 45	C9orf45	81571	NP_110441	NM_030814	4D	223522_at
Rasassociated protein Rap1	RBJ	51277	NP_057628	NM_016544	4D	223504 at
hydroxyacylglutathione hydrolaselike	HAGHL	84264	NP_115680	NM_032304	4D	223485_at
translocase of inner mitochondrial membrane 8 homo	TIMM8B	26521	NP_036591	NM_012459	4D	223478_at
phosphatidylinositol glycan, class M	PIGM	93183	NP_660150	NM_145167	4D	223470_at
calcium/calmodulindependent protein kinase kinase	CAMKKI	84254	NP_115670	NM_032294	4D	223460_at
peroxisome proliferative activated receptor, alpha	PPARA	5465	NP_001001928	NM_001001928	4D	223437_at
chromosome 7 open reading frame 36	C7orf36	57002	NP_064577	NM_020192	4D	223433 at

potassium voltagegated channel, subfamily H (eagre G proteincoupled receptor 81 G proteincoupled receptor 87 Calcium channel, voltagedependent, gamma subunit 7						
potassium voltagegated channel, subfamily H (eagre G proteincoupled receptor 81	CACNG7	59284	NP_114102	NM_031896	4D	224137_at
potassium voltagegated channel, subfamily H (eagre	GPR81	27198	NP_115943	NM_032554	4D	224131_at
nippocampus abundant gene transcriptiike 2	KCNH7	90134	NP_150375	NM_033272	4Đ	224099_at
	HIATL2	84278	NP_115694	NM_032318	40	224078 at
visual system homeobox 1 homolog, CHX10like (zebra	VSXI	30813	NP_955457	NM_199425	4D	224074_at
	ZCWPWI	55063	NP_060454	NM_017984	4D	223992 x at
	FLJ20859	64745	NP_073571	NM_022734	4D	223988 x at
IGFII mRNAbinding protein 2	IMP2	10644	NP_006539	NM_006548	4D	223963 s at
transmembrane protease, serine 3	TMPRSS3	64699	NP_115781	NM_032405	4D	223949 at
transmembrane protease, serine 3	TMPRSS3	64699	NP_076927	NM_024022	4D	223948 s_at
carbohydrate (Nacetylglucosamine 6O) sulfotransfer	CHST5	23563	NP_036258	NM_012126	ŧ	223942_x_at
3 solute carrier family 39 (zinc transporter), membe	SLC39A3	29985	NP_653165	NM_144564	4D	223917 s at
BCL6 corepressor	BCOR	54880	NP_065977	NM_020926	4D	223915_at
epsin 3	EPN3	55040	NP_060427	NM_017957	4D	223895 s_at
G proteincoupled receptor 132	GPR 132	29933	NP_037477	NM_013345	4D	223887_at
	C20orf24	55969	NP_061328	NM_018840	4D	223880_x_at
armadillo repeat containing 2	ARMC2	84071	NP_115507	NM_032131	4D	223866_at
	LOC51234	51234	NP_057538	NM_016454	4D	223857_x_at
orthopedia homolog (Drosophila)	OTP	23440	NP_115485	NM_032109	4D	223835_x_at
.G1 programmed cell death 1 ligand 1	PDCD1LG	29126	NP_054862	NM_014143	4D	223834_at
	KIAA1160	57461	NP_065752	NM_020701	4D	223831_x_at
	FLJ10052	55061	NP_060452	NM_017982	4D	223821_s_at
regulator of Gprotein signalling 18	RGS18	64407	NP_570138	NM_130782	4D	223809_at
GTP binding protein 2	GTPBP2	54676	NP_061969	NM_019096	4D	223789 s at
kinesin family member 9	KIF9	64147	NP_071737	NM_022342	4D	223778_at
dystrobrevin binding protein 1	DTNBPI	84062	NP_898861	NM_183040	4D	223763_at
	GTF2H2	2966	NP_001506	NM_001515	4D	223758 s at
	MGC13057	84281	NP_115697	NM_032321	4D	223754_at
	CIQTNF2	114898	NP_114114	NM_031908	4D	223749_at
mitochondrial ribosomal protein L4	MRPL4	51073	NP_057040	NM_015956	4D	223743 s_at

			-			-
thioredoxinlike 5	TXNL5	84817	NP 116120	NM 032731	4D	224511 s at
hypothetical protein MGC10744	MGC10744	84314	NP_898888	NM_183065	4D	224496 s_at
hypothetical protein MGC10744	MGC10744	84314	NP_898888	NM_183065	4D	224495_at
HECT domain containing 1	HECTDI	25831	NP_056197	NM_015382	4D	224481_s_at
leucine zipper, putative tumor suppressor 2	LZTS2	84445	NP_115805	NM_032429	4D	224473_x_at
PYM protein	PYM	84305	NP_115721	NM_032345	4D	224465 s at
chromosome 9 open reading frame 125	C9orf125	84302	NP_115718	NM_032342	4D	224458_at
hypothetical protein MGC12966	MGC12966	84792	NP_116095	NM_032706	4D	224452 s at
RIO kinase 1 (yeast)	RIOKI	83732	NP_113668	NM_031480	4D	224450 s at
ring finger protein 7	RNF7	9616	NP_899060	NM_183237	4D	224439 x at
chromosome 10 open reading frame 58	C10orf58	84293	NP_115709	NM_032333	4D	224435_at
mitochondrial translation optimization 1 homolog	MTO1	25821	NP_036255	NM_012123	4D	224430_s_at
mediator of RNA polymerase II transcription, subun	MED28	80306	NP_079481	NM_025205	4D	224416_s_at
histidine triad nucleotide binding protein 2	HINT2	84681	NP_115982	NM_032593	4D	224415_s_at
ring finger protein 7	RNF7	9616	NP_055060	NM_014245	4D	224395_s_at
ring finger protein 17	RNF17	56163	NP_114383	NM_031994	4D	224383_at
TAF7like RNA polymerase II, TATA box binding prote	TAF7L	54457	NP_079161	NM_024885	4D	224380_s_at
RAB18, member RAS oncogene family	RAB18		NP_067075	NM_021252	4D	224377_s_at
NDRG family member 3	NDRG3	57446	NP_114402	NM_032013	4D	224368 s_at
ubiquitinconjugating enzyme E2, J2 (UBC6 homolog,	UBE2J2	118424	NP_919297	NM_194316	4D	224347_x_at
mitochondrial ribosomal protein L51	MRPL51	51258	NP_057581	NM_016497	4D	224334 s at
mitochondrial ribosomal protein L36	MRPL36	64979	NP_115868	NM_032479	4D	224331 s at
mitochondrial ribosomal protein L27	MRPL27	51264	NP_683412	NM_148571	4D	224330_s_at
KIAA1287 protein	KIAA1287	57508	NP_065799	NM 020748	4D	224308 s_at
SECIS binding protein 2	SECISBP2	79048	NP_076982	NM_024077	4D	224250_s_at
Fas (TNFRSF6) associated factor 1	FAFI	11124	NP_008982	NM_007051	4D	224217_s_at
CGI30 protein	CGI30	51611	NP_057042	NM_015958	4D	224196_x_at
heat shock 70kDa protein 8	HSPA8	3312	NP 006588	NM_006597	4D	224187_x_at
chromosome X open reading frame 26	CXorf26	51260	NP_057584	NM 016500	4D	224177 s at
tripartite motifcontaining 4	TRIM4	89122	NP_148977	NM_033017	4D	224159_x_at
interleukin 17 receptor B	IL17RB	55540	NP_061195	NM_018725	4D	224156 x at

224516 s at	4D	NM_016463	NP 057547	51523	CXXC5	CXXC finger 5
224550 s_at	4D	NM_006069	NP_006060	10335	MRVII	murine retrovirus integration site 1 homolog
224563_at	4D	NM_006990	NP 008921	10163	WASF2	WAS protein family, member 2
224564_s_at	4D	NM_201428	NP_958831	10313	RTN3	reticulon 3
224575_at	4D	NM_018462	NP_060932	55845	C3orf10	chromosome 3 open reading frame 10
224584 at	4D	NM_014145	NP_054864	29058	C20orf30	chromosome 20 open reading frame 30
224593_at	4D	NM_152437	NP_689650	144348	ZFOC1	zinc finger protein ZFOC1
224595_at	4D	NM_080546	NP_536856	23446	CDW92	CDW92 antigen
224609_at	4D	NM_020428	NP_065161	57153	CTL2	CTL2 gene
224614_at	4D	NM_006141	NP_006132	1783	DNCL12	dynein, cytoplasmic, light intermediate polypeptid
224625_x_at	4D	NM_005770	NP_005761	10169	SERF2	small EDRKrich factor 2
224632_at	4D	NM_182679	NP_872620	54865	GPATC4	G patch domain containing 4
224634_at	4D	NM_015590	NP_056405	54865	GPATC4	G patch domain containing 4
224637_at	4D	XM_376043	XP_376043	400948	LOC400948	similar to RIKEN cDNA 2310016E02
224640 at	4D	NM_139015	NP_620584	121665	SPPL3	signal peptide peptidase 3
224655_at	4D	NM_016282	NP_057366	50808	AK3L1	adenylate kinase 3 like 1
224657_at	4D	NM_018948	NP_061821	54206	MIG6	mitogeninducible gene 6
224661_at	4D	NM_032906	NP_116295	84992	MGC14156	hypothetical protein MGC14156
224662 at	4D	NM_004521	NP_004512	3799	KIF5B	kinesin family member 5B
224669_at	4D	NM_033542	NP_291020	55861	C20orf35	chromosome 20 open reading frame 35
224713_at	4D	NM_032390	NP_115766	84365	MK167IP	MK167 (FHA domain) interacting nucleolar phosphopr
224719_s_at	4D	NM_138425	NP_612434	113246	GRCC10	likely ortholog of mouse gene rich cluster, C10 ge
224721_at	4D	NM_032168	NP_115544	84128	FLJ12519	hypothetical protein FLJ12519
224736_at	4D	NM_018237	NP_060707	55749	CCARI	cell division cycle and apoptosis regulator 1
224739_at	4D	NM_001001852	NP_001001852	415116	PIM3	pim3 oncogene
224768_at	4D	NM_017969	NP_060439	55677	FLJ10006	hypothetical protein FLJ10006
224779 s_at	4D	NM_032231	NP_115607	84191	FLJ22875	hypothetical protein FLJ22875
224780_at	4D	NM_032905	NP_116294	84991	RBM17	RNA binding motif protein 17
224785 at	4D	NM_182565	NP_872371	283991	MGC29814	hypothetical protein MGC29814
224787 s at	4D	NM_021252	NP_067075	22931	RAB18	RAB18, member RAS oncogene family
224797_at	4D	NM_020801	NP_065852	57561	ARRDC3	arrestin domain containing 3
224812_at	4D	NM_152740	NP_689953	11112	HIBADH	3hydroxyisobutyrate dehydrogenase
224819_at	4D	NM 153333	NP 699164	90843	TCEAL8	transcription elongation factor A (SII)like 8

G elongation factor, mitochondrial 1	GFM1	85476	NP 079272	NM 024996	4D	225153_at
KIAA1718 protein	KIAA1718	80853	XP_376680	XM_376680	4D	225142_at
Kruppellike factor 3 (basic)	KLF3	51274	NP_057615	NM_016531	4D	225133_at
rabaptin, RAB GTPase binding effector protein 1	RABEP1	9135	NP_004694	NM_004703	4D	225092_at
synovial apoptosis inhibitor 1, synoviolin	SYVNI	84447	NP_115807	NM_032431	4D	225090_at
hypothetical protein FLJ31153	FLJ31153	123811	NP_653201	NM_144600	4D	225088_at
cleavage and polyadenylation specific factor 3, 73	CPSF3	51692	NP_057291	NM_016207	4D	225082_at
RAB2B, member RAS oncogene family	RAB2B	84932	NP_116235	NM_032846	4D	225074_at
chromosome 6 open reading frame 68	C6orf68	116150	NP_612468	NM_138459	4D	225071_at
DnaJ (Hsp40) homolog, subfamily A, member 4	DNAJA4	55466	NP_061072	NM_018602	4D	225061_at
angiotensin II receptorassociated protein	AGTRAP	57085	NP_065083	NM_020350	4D	225059_at
zinc finger protein 512	ZNF512	84450	NP_115810	NM_032434	4D	225050_at
solute carrier family 15, member 4	SLC15A4	121260	NP_663623	NM_145648	4D	225043_at
chromosome 9 open reading frame 105	C9orf105	401505	NP_001001790	NM_001001790	4D	225036_at
chromodomain helicase DNA binding protein 6	CHD6	84181	NP_115597	NM_032221	4D	225031_at
hypothetical gene supported by BC032431	LOC389203	389203	XP_371691	XM_371691	4D	225014_at
ataxin 7like 3	ATXN7L3	56970	XP_375456	XM_375456	4D	224969_at
solute carrier family 26 (sulfate transporter), me	SLC26A2	1836	NP_000103	NM_000112	4D	224963_at
solute carrier family 26 (sulfate transporter), me	SLC26A2	1836	NP_000103	NM_000112	4D	224959_at
82kD FMRP Interacting Protein	182FIP	57532	NP_065823	NM_020772	4D	224956_at
mitochondrial ribosomal protein S24	MRPS24	64951	NP_114403	NM_032014	4D	224948_at
chromosome 22 open reading frame 16	C22orf16	400916	NP_998885	NM_213720	4D	224932_at
myeloidassociated differentiation marker	MYADM	91663	NP_612382	NM_138373	4D	224920_x_at
hom	1 TIVILIVI I	72007	141 001001303	MM 001001303	ė	224713_3_at
chromosome 9 open reading frame 123	C9ort123	90871	XP_372075	XM_372075	4	224879 at
chromosome 9 open reading frame 123	C9orf123	90871	XP_372075	XM_372075	4D	224860 at
tumor protein p53 inducible nuclear protein 2	TP53INP2	58476	NP_067025	NM 021202	4D	224836 at
cytoplasmic polyadenylation element binding protei	CPEB4	80315	NP_085130	NM_030627	4D	224831_at
cytoplasmic polyadenylation element binding protei	CPEB4	80315	NP_085130	NM_030627	ŧ	224828_at
family with sequence similarity 36, member A	FAM36A	116228	NP_932342	NM_198076	4D	224824_at
family with sequence similarity 36, member A	FAM36A	116228	NP_932342	NM_1980/6	40	224820 at

chromosome 1 open reading frame 19	Clorf19	116461	NP_443197	NM_052965	4D	225399_at
dedicator of cytokinesis 7	DOCK7	85440	NP_212132	NM_033407	4D	225384_at
endothelial cell adhesion molecule	ESAM	90952	NP_620411	NM_138961	4D	225369 at
hypothetical protein PP2447	PP2447	80305	NP_079480	NM_025204	4D	225360_at
homolog of yeast TIM14	TIM14	131118	NP_957713	NM_201261	4D	225359_at
chromosome 3 open reading frame 6	C3orf6	152137	NP_848018	NM_178335	4D	225331_at
acylCoenzyme A binding domain containing 6	ACBD6	84320	NP_115736	NM_032360	4D	225317_at
COMM domain containing 6	COMMD6	170622	NP_987091	NM_203495	4D	225312_at
polyribonucleotide nucleotidyltransferase 1	PNPT1	87178	NP_149100	NM_033109	4D	225291_at
hypothetical protein AL133206	LOC64744	64744	NP_073570	NM_022733	4D	225282_at
spermidine/spermine N1acetyltransferase 2	SAT2	112483	NP_597998	NM_133491	4D	225272_at
THHike (Drosophila)	THIL	51497	NP_945327	NM_198976	4D	225261 x at
mitochondrial ribosomal protein L32	MRPL32	64983	NP_114109	NM_031903	4D	225260_s_at
mitochondrial ribosomal protein L35	MRPL35	51318	NP_663619	NM_145644	4D	225255_at
sarcolemma associated protein	SLMAP	7871	NP_009090	NM_007159	4D	225243_s_at
musashi homolog 2 (Drosophila)	MSI2	124540	NP_620412	NM_138962	4D	225237_s_at
hypothetical LOC83640	MGC2560	83640	NP_113640	NM_031452	4D	225210_s_at
mitochondrial ribosomal protein L14	MRPL14	64928	NP_115487	NM_032111	4D	225201 s_at
mitochondrial ribosomal protein S26	MRPS26	64949	NP_110438	NM_030811	4D	225196_s_at
zinc finger, CSL domain containing 2	ZCSL2	285381	NP_996662	NM_206831	4D	225195_at
Ras association (RalGDS/AF6) and pleckstrin homolo	RAPHI	65059	NP_998754	NM_213589	4D	225189_s_at
homolo			1	1		1
Ras association (RalGDS/AF6) and pleckstrin	RAPHI	65059	NP 998754	NM 213589	4D	225188 at
PRO0149 protein	PRO0149	29035	NP_054836	NM 014117	4D	225183_at
chromosome 21 open reading frame 4	C21orf4	757	NP_006125	NM_006134	4D	225182_at
tetratricopeptide repeat domain 14	TTC14	151613	NP_597719	NM_133462	4D	225180_at
RAB11 family interacting protein I (class I)	RAB11FIP1	80223	NP_001002814	NM_001002814	4D	225177_at
Rho GTPase activating protein 18	ARHGAP18	93663	NP_277050	NM_033515	4D	225173_at
protein phosphatase 1, regulatory (inhibitor) subu	PPPIRIB	84152	NP_852606	NM_181505	4D	225165_at
FERM domain containing 4A	FRMD4A	55691	NP_060497	NM_018027	4D	225163_at
hypothetical protein MGC4549	MGC4549	84337	NP_115753	NM_032377	4D	225156_at
synapse associated protein 1, SAP47 homolog (Droso	SYAPI	94056	NP_116185	NM_032796	4D	225154_at

methionine sulfoxide reductase B3	MSRB3	253827	NP 932346	NM 198080	40	225782 at
chromosome 6 open reading frame 93	C6orf93	84946	NP_116249	NM_032860	40	225748_at
zinc finger, DHHC domain containing 8	ZDHHC8	29801	NP_037505	NM_013373	4D	225744 at
RNA pseudouridylate synthase domain containing 3	RPUSD3	285367	NP_775930	NM_173659	4D	225743_at
THUMP domain containing 3	THUMPD3	25917	NP 056268	NM_015453	4D	225741 at
mitochondrial ribosomal protein L55	MRPL55	128308	NP_852129	NM 181464	4D	225719 s at
KIAA1715	KIAA1715	80856	NP_085153	NM_030650	4D	225718_at
hypothetical protein BC009518	LOC90799	90799	NP_612372	NM_138363	4D	225705_at
TIGAI	TIGAI	114915	NP_443726	NM_053000	4D	225698_at
family with sequence similarity 33, member A	FAM33A	348235	NP_872426	NM_182620	4D	225686_at
interferon (alpha, beta and omega) receptor 1	IFNAR1	3454	NP_000620	NM_000629	4D	225661_at
hypothetical protein LOC339745	LOC339745	339745	NP_001001664	NM_001001664	4D	225659_at
ubiquitinconjugating enzyme E2E 2 (UBC4/5 homolog,	UBE2E2	7325	NP_689866	NM_152653	\$	225651_at
similar to hypothetical protein 9530023G02	MGC90512	121642	NP_001001655	NM_001001655	4D	225625_at
UDPGlcNAc:betaGal beta1,3Nacetylglucosaminyltransf	B3GNT5	84002	NP_114436	NM_032047	4D	225612_s_at
hypothetical protein FLJ31795	FLJ31795	124808	NP_653210	NM_144609	40	225607_at
chromosome 9 open reading frame 19	C9orf19	152007	NP_071738	NM_022343	4D	225604_s_at
SH3 multiple domains 2	SH3MD2	57630	NP_065921	NM_020870	4D	225589_at
putative acylCoA dehydrogenase	FLJ12592	84129	NP_115545	NM_032169	4D	225573_at
chromosome 3 open reading frame 19	C3orf19	51244	NP_057558	NM_016474	4D	225559_at
zinc finger, CCHC domain containing 9	ZCCHC9	84240	NP_115656	NM_032280	4D	225538_at
translocase of inner mitochondrial membrane 23 hom	TIMM23	10431	NP_006318	NM_006327	4D	225535_s_at
gamma			1	-		
CCAAT/enhancer binding protein (C/EBP).	CEBPG	1054	NP 001797	NM 001806	4Đ	225527 at
chromosome 20 open reading frame 81	C20orf81	64773	NP_073597	NM_022760	4D	225505_s_at
transmembrane protein 18	TMEM18	129787	NP_690047	NM_152834	4D	225489_at
mesoderm induction early response 1	MIERI	57708	NP 065999	NM_020948	4D	225475_at
nucleoporin 35kDa	NUP35	129401	NP_612142	NM_138285	4D	225470_at
glycerol3phosphate acyltransferase, mitochondrial	GPAM	57678	NP_065969	NM_020918	4D	225424_at
glycerol3phosphate acyltransferase, mitochondrial	GPAM	57678	NP_065969	NM_020918	4D	225420_at
ring ringer protein 149	RNF149	284996	NP_775918	NM_173647	4D	225414_at

	100285626	285636	NP 787117	NM 175921	4D	226159 at
retinoic acid induced 1	RAII	10743	NP_109590	NM_030665	4D	226143 at
hypothetical protein MGC23280	MGC23280	147015	NP_653284	NM_144683	4D	226121_at
TRAFinteracting protein with a forkheadassociated	TIFA	92610	NP_443096	NM_052864	40	226117_at
fibronectin type 111 domain containing 5	FNDC5	252995	NP 715637	NM_153756	4	226097 at
membrane protein, palmitoylated 5 (MAGUK p55 subfa	MPP5	64398	NP_071919	NM_022474	ŧ	226092_at
synaptotagmin XIII	SYT13	57586	NP_065877	NM_020826	4	226086 at
caspase 2, apoptosisrelated cysteine protease (neu	CASP2	835	NP_116764	NM_032982	40	226036_x_at
disrupted in renal carcinoma 2	DIRC2	84925	NP_116228	NM_032839	40	226026_at
copper metabolism (Murr1) domain containing	COMMDI	150684	NP_689729	NM_152516	4D	226024_at
hypothetical protein Ells l	Ells1	222166	NP_690006	NM_152793	4D	226018_at
coiledcoil domain containing 12	CCDC12	151903	NP_653317	NM_144716	40	226011_at
HESB like domain containing 1	HBLD1	122961	NP_919255	NM_194279	4D	226007_at
cleavage and polyadenylation specific factor 2, 10	CPSF2	53981	NP_059133	NM_017437	Đ	225994_at
hypothetical protein MGC15677	MGC15677	84964	NP_116267	NM_032878	4B	225969 at
adult retina protein	LOC153222	153222	NP_705835	NM_153607	4	225957_at
zinc finger protein 655	ZNF655	79027	NP_612503	NM_138494	ŧ	225945_at
CREBBP/EP300 inhibitor 2	CR12	163126	NP_694964	NM_153232	4D	225936_at
FLJ45445 protein	FLJ45445	399844	NP_001004321	NM_001004321	4D	225899 x_at
hypothetical protein FLJ13089	FLJ13089	81008	NP_079229	NM_024953	Đ	225888_at
solute carrier family 35, member B4	SLC35B4	84912	NP_116215	NM_032826	4D	225882_at
solute carrier family 35, member B4	SLC35B4	84912	NP_116215	NM_032826	4D	225881_at
hypothetical protein dJ462O23.2	DJ462O23.2	57185	NP_065181	NM_020448	4D	225876_at
solute carrier family 35, member F5	SLC35F5	80255	NP_079457	NM_025181	4D	225872_at
glucosaminephosphate Nacetyltransferase 1	GNPNAT1	64841	NP_932332	NM_198066	4D	225853 at
ankyrin repeat domain 17	ANKRD17	26057	NP_942592	NM_198889	4D	225852_at
chromosome 6 open reading frame 83	C6orf83	113402	NP_660152	NM_145169	4D	225850 at
chromosome 6 open reading frame 83	C6orf83	113402	NP_660152	NM_145169	4D	225849 s at
hypothetical protein MGC13204	MGC13204	83695	NP_113653	NM_031465	4D	225837_at
hypothetical protein MGC13204	MGC13204	83695	NP_113653	NM_031465	4D	225836 s at
hypothetical gene supported by AL449243	LOC91689	91689	NP 201575	NM_033318	4D	225795_at
methionine sulfoxide reductase B3	MISKBS	233027	1AL 2020	1100000		

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226614_s_at 4D	226611 s at 4D	226603_at 4D	226564 at 4D	226537 at 4D	226531_at 4D	226524_at 4D	226515_at 4D	226507_at 4D	226489_at 4D	226487_at 4D	226470_at 4D	226443_at 4D	226442_at 4D	226422_at 4D	226394_at 4D	226385_s_at 4D	226378_s_at 4D	226339_at 4D	226338_at 4D	226321_at 4D	226319_s_at 4D	226301_at 4D	226296_s_at 4D	226276_at 4D	226267_at 4D	226239_at 4D	226194 at 4D	226177_at 4D	226175_at 4D	220109 at 4D
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NM_053279	NM_181716	NM_152703	NM_020863	NM_138571	NM_032790	NM_173824	NM_145265	NM_002576	NM_020698	NM_032829	NM_052830	NM_138333	NM_032548	NM_016570	NM_017824	NM_138446	NM_152482	NM_139169	NM_018710	XM_371760	NM_005782	NM_052831	NM_031280	NM_174909	NM_130469	NM_153342	NM_032436	NM_016433	NM_173810	7000C ININI
NP 444509	NP_859067	NP_689916	NP_065914	NP_612638	NP_116179	NP_776185	NP_660308	NP_002567	NP_065749	NP_116218	NP_439895	NP_612206	NP_115937	NP_057654	NP_060294	NP_612455	NP_689695	NP_631908	NP_061180	XP_371760	NP_005773	NP_439896	NP_112570	NP_777569	NP_569736	NP_699173	NP_115812	NP_057517	NP_776171	
83648	201161	219285	57623	135114	84876	285237	133957	5058	57458	84915	2686	116224	80325	51290	54708	115416	148223	142940	55529	116068	10189	116843	64960	153339	122953	129303	283489	51228	283237	
C8orf13	PRR6	C7orf6	ZNF406	HINT3	FLJ14466	MGC26717	LOC133957	PAKI	TMCC3	FLJ14721	GGTL3	C9orf42	ABTB1	PTXI	MARCH5	C7orf30	C19orf25	TRUBI	DKFZp762O076	LOC116068	THOC4	C6orf192	MRPS15	MGC23909	JDP2	FLJ90024	C13orf8	GLTP	MGC29649	0.00
chromosome 8 open reading frame 13	proline rich 6	chromosome 7 open reading frame 6	zinc finger protein 406	histidine triad nucleotide binding protein 3	hypothetical protein FLJ14466	hypothetical protein MGC26717	similar to RIKEN cDNA 0610011N22	p21/Cdc42/Rac1activated kinase 1 (STE20 homolog, y	transmembrane and coiledcoil domains 3	hypothetical protein FLJ14721	gammaglutamyltransferaselike 3	chromosome 9 open reading frame 42	ankyrin repeat and BTB (POZ) domain containing	PTX1 protein	membraneassociated ring finger (C3HC4) 5	chromosome 7 open reading frame 30	chromosome 19 open reading frame 25	TruB pseudouridine (psi) synthase homolog 1 (E. co	hypothetical protein DKFZp762O076	hypothetical protein LOC116068	THO complex 4	chromosome 6 open reading frame 192	mitochondrial ribosomal protein S15	hypothetical protein MGC23909	jun dimerization protein 2	fastinginducible integral membrane protein TM6P1	chromosome 13 open reading frame 8	glycolipid transfer protein	hypothetical protein MGC29649	one omening mercer at

4D NM 025240 NP 025359 29843 SUP ILDI 4D NM 025240 NP 002633 5337 PLDI 4D NM 025240 NP 116129 84825 MCC3391 4D NM 032740 NP 116129 84825 MCC3391 4D NM 032740 NP 003632 10491 CRTAP 4D NM 13236 NP 006362 10491 CRTAP 4D NM 173826 NP 776187 28343 DKT2p313N062 4D NM 173826 NP 775187 28343 DKT2p313N062 4D NM 173826 NP 775187 28343 DKT2p313N062 4D NM 173826 NP 872578 64962 KIAA1991 4D NM 173826 NP 872578 64962 KIAA1991 4D NM 187833 NP 849157 152483 DKT2p313N062 4D NM 197964 NP 932068 154791 HSPC268 4D NM 197964 NP 932068 154791 HSPC268 4D NM 197964 NP 937790 16236 TWSTRB 4D NM 198147 NP 937790 16236 COC116236 4D NM 182493 NP 8937791 119587 CKNAP5 4D NM 182493 NP 937790 16236 CCC116236 4D NM 182493 NP 937790 16236 CCC116236 4D NM 182493 NP 937790 16236 GRED1 4D NM 182493 NP 937891 19818 DOCK11 4D NM 152407 NP 653229 139818 DOCK11 4D NM 152407 NP 683229 139818 DOCK11 4D NM 152407 NP 683229 139818 DOCK11 4D NM 152407 NP 683259 139818 DOCK11 4D NM 152407	hypothetical protein MGC40107	MGC40107	254863	NP_689979		4D	227063_at
MN 012524 NP 025356 298.3 SUP	protein kinase C and case	PACSIN1	29993	NP_065855	NM_020804		227053_at
4D NM 032540 NP 035369 29843 SUP ILDI 4D NM 032740 NP 116129 84825 MCG391 4D NM 032740 NP 116129 84825 MCG391 4D NM 032740 NP 116129 84826 MCG391 4D NM 14526 NP 006362 10491 CRYAP 4D NM 173826 NP 776187 28313 DKF2p313N062 4D NM 173826 NP 776187 28313 DKF2p313N062 4D NM 173826 NP 776187 28323 DKF2p313N062 4D NM 173826 NP 975897 284129 SLC26A11 4D NM 173826 NP 849157 152485 LOC(152485 NP 849157) 4D NM 173826 NP 932068 154791 HSPC268 4D NM 198143 NP 932068 154991 HSPC268 4D NM 198144 NP 640337 15495 FAHDI 4D NM 19244 NP 640337 116236 DCC(16236 NP 93790 116236 NP 93790 116236 STAPPS 4D NM 19244 NP 640337 119587 CC(16236 NP 93790 116236 STAPPS 4D NM 19244 NP 640337 119587 CC(16236 NP 901001936 S4632 KIAA1914 NP 648249 SP706 S4632 SP804 SP8	chromosome 14 open rea-	C14orf24	283635	NP_775878	NM_173607		227029 at
4D NM 02562 NP 025369 29843 SNP1 4D NM 02562 NP 02633 5337 PLDI 4D NM 02562 NP 02633 5337 PLDI 4D NM 025740 NP 116159 84826 MGC3391 4D NM 032740 NP 116159 84826 MGC3391 4D NM 145266 NP 776187 285343 DKF25313N062 4D NM 173825 NP 776187 285343 DKF25313N062 4D NM 173826 NP 776187 285343 DKF25313N062 4D NM 173825 NP 575187 285343 DKF25313N062 4D NM 173825 NP 549157 152485 LOC152485 4D NM 19764 NP 64925 152485 LOC152485 4D NM 19764 NP 63037 134957 STYSBFS 4D NM 19764 NP 640337 134957 CPXA276 4D NM 198147 NP 640337 134957 CPXA276 4D NM 198148 NP 637790 116236 LOC116236 4D NM 198148 NP 937791 119587 CPXA276 4D NM 182493 NP 689807 116342 KIAA1914 4D NM 182349 NP 689807 116342 KIAA1914 4D NM 182349 NP 689807 116342 KIAA1914 4D NM 132383 NP 16274 84971 APG4D 4D NM 132392 NP 689493 DAST6 ACPL2 4D NM 132393 NP 689493 DAST6 ACPL2 4D NM 15284 NP 69679 DAST6 ACPL2 4D NM 15284 NP 69679 AGP43 BCAP12	U11/U12 snRNP 65K	RNP	55599	NP_060089	NM_017619		226975 at
4D NM 014554 NP 085369 29843 ENPI 4D NM 014554 NP 065363 5337 PLDI 4D NM 032740 NP 016129 844826 MGC5391 NM 0145266 NP 606399 134492 NUDCD2 4D NM 032740 NP 175897 294129 SLC25A11 NM 096371 NP 065362 164491 CRT.AP 4D NM 173826 NP 175897 294129 SLC25A11 NM 178336 NP 175897 152485 LOC152485 NM 178338 NP 189157 152485 LOC152485 NM 178341 NM 178344 NP 640337 154537 174151NB NM 198144 NP 640337 154536 LOC1 16236 NM 152493 NM 152594 NP 640379 116236 LOC1 16236 NM 152594 NP 659807 161742 SPREDI NM 152582 NP 659807 154286 GRPEL2 NM 152582 NP 659803 15426 GRPEL2 NM 152582 NP 659639 15426 GRPEL2 NM 152582 NP 166974 85403 EAFT 1400 NM 152582 NP 659639 15426 GRPEL2 NM 152603 NP 1540974 85403 EAFT 1400 NM 152582 NP 169074 85403 EAFT 1400 NM 152682 NP 169074 85403 EAFT 1400 NM 152682 NP 169074 85403 EAFT 1400 NM 152603 NP 156003 400924 LOC100924	ralA binding protein 1	RALBPI	10928	NP 006779	NM_006788		226957 x at
4D NM 014524 NP 055369 23943 EAPH 1 4D NM 002562 NP 002563 5337 PLDI 1 4D NM 002562 NP 0605369 134492 MDCC391 NM 015256 NP 276187 284129 SLC26A11 NM 125266 NP 276187 285343 DKF251310662 NP 176187 28546 DKF25131062 NP 176187 28546 DKF2513106 NP 176187 28546 DKF2513	hypothetical gene supports	LOC400924	400924	XP_376003			226956_at
4D NM 014554 NP 035369 2943 SIRPE 4D NM 025240 NP 026533 S337 PLDI 4D NM 145266 NP 660309 SIRPE 4D NM 173626 NP 660309 SIRPE 4D NM 173626 NP 775897 294129 SLC26A11 4D NM 173626 NP 175897 29513 SNF2D313N062 4D NM 173626 NP 872578 64945 MRE99 4D NM 173626 NP 112485 SIRS9 FAHDI 4D NM 173626 NP 600337 SIRSP	ELL associated factor I	EAFI	85403	NP_149074	NM_033083		226952_at
4D NM 014554 NP 025369 29843 ENPI 4D NM 02562 NP 025359 5337 PLDI 4D NM 032740 NP 116129 84826 MGC5391 AD NM 032740 NP 116129 124492 CRT.p 4D NM 032761 NP 065369 1134492 SUC26A11 AD NM 173266 NP 775897 294129 SUC26A11 AD NM 173266 NP 775897 294129 SUC26A11 AD NM 173266 NP 175897 294129 SUC26A11 AD NM 173268 NP 175897 294129 SUC26A11 AD NM 178236 NP 91578 254235 KIAA1991 AD NM 178238 NP 89157 152485 COC152485 AD NM 178238 NP 89157 152485 LOC152485 AD NM 197964 NP 01002926 21839 FAHD1 AD NM 198144 NP 640337 114926 FL123476 AD NM 198144 NP 649376 116236 LOC1625 AD NM 198148 NP 01001956 84622 KIAA1914 AD NM 152549 NP 00001956 RIG22 KIAA1914	acid phosphataselike 2	ACPL2	92370	NP_689495	NM_152282		226925 at
4D NM 014554 NP 055369 29843 SEDIP 4D NM 02562 NP 062539 5337 PLDI 4D NM 032740 NP 116129 84826 MGC5391 4D NM 032740 NP 0605399 134492 NUDCD2 4D NM 135266 NP 776187 28313 DKP25131062 4D NM 178326 NP 176187 28313 DKP25131062 4D NM 182640 NP 872578 285343 DKP25131062 4D NM 178836 NP 89157 152485 LOC152485 4D NM 178838 NP 112485 81889 FAHD1 4D NM 197964 NP 032068 154791 HSPC268 4D NM 197964 NP 937790 16236 LOC16236 4D NM 198248 NP 937790 16236 LOC16236 4D NM 198249 NP 937790 16236 LOC16236 4D NM 198147 NP 937790 16236 LOC16236 4D NM 198148 NP 01001956 R4632 KIAA 1914 4D NM 032888 NP 116274 SPREDI	GrpElike 2, mitochondrial (GRPEL2	134266	NP_689620	NM_152407		226881_at
4D NM 014554 NP 035369 2984 SH899 LC25A11 4D NM 025240 NP 060359 3337 PLDI 4D NM 145266 NP 660369 134492 CRTAP 4D NM 173626 NP 660369 134492 CRTAP 4D NM 173626 NP 775897 284129 SLC26A11 4D NM 173626 NP 775897 284129 SLC26A11 4D NM 173626 NP 775897 284129 SLC26A11 4D NM 173626 NP 175897 285343 DKF2p313N662 4D NM 182640 NP 872578 64945 MRE99 4D NM 178833 NP 849157 152485 LCC152485 4D NM 01000226 NP 112485 81889 FAHDI 4D NM 139944 NP 600337 134957 STXBPS 4D NM 139944 NP 600337 134957 STXBPS 4D NM 1398147 NP 689706 149076 LCC16236 4D NM 198147 NP 937790 116236 LCC116236 4D NM 198148 NP 937790 119587 CPXXI2 4D NM 198148 NP 937790 149076 CPXXI2 4D NM 152594 NP 689807 161742 SPREDI 4D NM 152594 NP 689807 161742 SPREDI 4D NM 152885 NP 16574 84971 APG-4D	dedicator of cytokinesis 11	DOCKII	139818	NP_653259	NM_144658		226875_at
4D NM 032740 NP 062539 S337 PLDI 4D NM 032740 NP 062539 S337 PLDI 4D NM 032740 NP 16129 S4826 MGC5391 AD NM 032740 NP 16129 S4826 MGC5391 AD NM 145266 NP 603509 134492 CRT,AP AD NM 173262 NP 775897 284129 SLC25A11 AD NM 173262 NP 775897 284129 SLC25A11 AD NM 173262 NP 776187 285343 DKF2/3131062 AD NM 173264 NP 776187 285343 DKF2/3131062 AD NM 173264 NP 915886 254225 K1AA1991 AD NM 173264 NP 915886 254225 K1AA1991 AD NM 173264 NP 9152578 64963 MRPS9 AD NM 173268 NP 819157 152485 LOC152485 AD NM 173264 NP 912485 S1889 FAHDI AD NM 197964 NP 912068 154791 HSPC28 AD NM 198147 NP 917790 116236 LOC16236 AD NM 198147 NP 937790 116236 LOC16236 AD NM 198148 NP 93791 116236 LOC16236 AD NM 198148 NP 93790 116236 LOC16236 AD NM 198148 NP 93791 116236 LOC16236 AD NM 198148 NP 93791 116236 LOC16236 AD NM 152493 NP 689706 1199076 FLU25476 AD NM 152493 NP 689706 1199076 PCPXA2 AD NM 152594 NP 689707 161742 SPRED1	APG4 autophagy 4 homolog	APG4D	84971	NP_116274	NM_032885		226871_s_at
AD	sproutyrelated, EVH1 domai	SPRED1	161742	NP_689807	NM_152594		226837_at
4D NM_014554 NP_035369 29843 SENPI 4D NM_002662 NP_002533 S337 PLDI 4D NM_103264 NP_116129 84826 MGC5391 4D NM_145266 NP_660309 134492 CRTAP 4D NM_173626 NP_776187 285129 SLC26A1.11 4D NM_173826 NP_776187 285129 SLC26A1.11 4D NM_173826 NP_776187 285129 SLC26A1.11 4D NM_173826 NP_776187 28513 DKF2p313N062 4D NM_173826 NP_879578 64945 MRE99 4D NM_178833 NP_849157 152485 LCC152485 4D NM_0102026 NP_112485 81889 FAHD1 4D NM_0102026 NP_600337 134957 STXBPS 4D NM_139244 NP_600337 134957 STXBPS 4D NM_139244 NP_600370 114026 LCC116236 4D NM_139244 NP_609790 116236 LCC116236 4D NM_139244 NP_639790 116236 LCC116236 4D NM_152493 NP_837791 119387 CPXXIZ	KIAA1914	KIAA1914	84632		NM_001001936		226829_at
AD	carboxypeptidase X (M14 far	CPXM2	119587	NP_937791	NM_198148		226824 at
4D NM 014554 WP 025369 29843 SENDI 4D NM 02562 NP 025363 5337 PLDI 4D NM 032740 NP_116129 84826 MGC5391 4D NM 0352740 NP_176187 284129 SLC26A11 4D NM 173826 NP_776187 285343 DKF251318062 4D NM 182546 NP_776187 285343 DKF251318062 4D NM 182640 NP_872578 C54225 K1AA1991 4D NM 178836 NP_89157 152485 LOC152485 4D NM 178836 NP_112485 81889 FAHD1 4D NM 197964 NP_01002926 221830 TWISTNB 4D NM 197964 NP_603997 154957 STABPS 4D NM 197964 NP_603997 154957 STABPS	FLJ25476 protein	FLJ25476	149076	NP_689706	NM_152493		226820_at
4D NM_014554 NP_065369 29843 SENPI 4D NM_002662 NP_065359 5337 PLDI 4D NM_002662 NP_0602653 5337 PLDI 4D NM_002740 NP_116129 84826 MGC5391 4D NM_06371 NP_066329 134492 NUDCD2 4D NM_173626 NP_776187 284129 SLC26A11 4D NM_173826 NP_776187 285343 DK72p313N062 4D NM_173826 NP_776187 285343 DK72p313N062 4D NM_182640 NP_872578 64965 MRPS9 4D NM_182640 NP_872578 64965 MRPS9 4D NM_031208 NP_112485 81889 FAHDI 4D NM_01002926 NP_030068 218481 HSPC268 4D NM_01002926 NP_01002926 21880 TWSTNBPS	hypothetical protein LOCI16	LOC116236	116236	NP_937790	NM_198147		226796_at
4D NM 014554 NP 085369 29843 SENPI 4D NM 02562 NP 065363 5337 PLDI 4D NM 032740 NP 116129 84826 MGC5391 4D NM 032740 NP 116129 134826 MGC5391 4D NM 145266 NP 660399 134492 NUDCD2 4D NM 173626 NP 775897 284129 SLC26A11 4D NM 173826 NP 775887 284129 SLC26A11 4D NM 173826 NP 776187 285343 DKF2β313N062 4D NM 182640 NP 872578 64965 MRPS9 4D NM 178833 NP 849157 152485 LOC153485 4D NM 031208 NP 1212485 81889 FAHDI 4D NM 197964 NP 932068 154791 HSPC268 4D NM 001002926 NP 901002926 221839 IWISTNB	syntaxin binding protein 5 (to	STXBP5	134957	NP_640337			226794_at
AD	TWIST neighbor	TWISTNB	221830	NP_001002926			226784_at
4D NM_01454 NP_05359 29843 SENPI 4D NM_01454 NP_05359 29843 SENPI 4D NM_052740 NP_16129 84826 MGC5391 4D NM_14526 NP_60399 154492 NUDCD2 4D NM_17326 NP_76187 284129 SLC26A11 4D NM_173826 NP_776187 285343 DKFZp313N062 4D NM_182640 NP_572578 64965 MRPS9 4D NM_18236 NP_872578 64965 MRPS9 4D NM_18236 NP_112485 81889 FAHD1	hypothetical protein HSPC26	HSPC268	154791	NP_932068	NM_197964		226780_s_at
4D NM 01454 NP 055569 29843 SENDI 4D NM 02562 NP 065539 3337 PLDI 4D NM 1632740 NP 116129 84825 MGC5391 4D NM 145266 NP 666399 134492 CRTAP 4D NM 173626 NP 775897 284129 SLC28A11 4D NM 173626 NP 775897 285133 DKF2p313N062 4D NM 173826 NP 879157 25485 LOC152485	fumarylacetoacetate hydrolase	FAHDI	81889		NM_031208		226767_s_at
4D NM_01454 NP_085369 2983 SENP1 4D NM_01454 NP_085369 2983 SENP1 4D NM_002662 NP_085363 5337 PLD1 4D NM_032740 NP_16129 84826 MGC5391 4D NM_145266 NP_606309 154492 NUDCD2 4D NM_173626 NP_76897 284129 SLC26A11 4D NM_173626 NP_776187 285343 DKFZp313N062 4D NM_182640 NP_872578 64965 NRPS9	hypothetical protein LOC152-	LOC152485	152485	NP_849157	NM_178835		226764 at
4D NM 01454 NP 085369 29843 SENDI 4D NM 00262 NP 005369 3337 PLDI 4D NM 032740 NP 116129 84826 MGC5391 4D NM 145266 NP 660399 134492 NUDCP29 4D NM 173626 NP 775897 284129 SLC26A11 4D NM 173826 NP 776187 285343 DKF2p313N062 4D NM 173826 NP 776187 285343 DKF2p313N062	mitochondrial ribosomal prote	MRPS9	64965	NP 872578	NM_182640		226749_at
4D NM 01454 NP 055369 29843 SEND1 4D NM 002662 NP 002536 5337 PLD1 4D NM 002740 NP 116129 84825 MIDC 023 4D NM 1032740 NP 606309 134492 MIDC 023 4D NM 006371 NP 006362 10491 CRT AP 4D NM 173626 NP 776187 283343 DKF Zp31310662	hypothetical protein KIAA199	K1AA1991	254225	XP_495886	XM_495886		226739_at
4D NM 01454 NP 085369 29843 SENDI 4D NM 002662 NP 002536 3337 PLDI 4D NM 032740 NP 116129 84826 MGC5391 4D NM 032770 NP 006562 116491 CRTAP 4D NM 173626 NP 775897 284129 SLCZ6A11	hypothetical protein DKFZp3	DKFZp313N062	285343	NP_776187	NM_173826		226688_at
4D NM_014534 NP_085369 29843 SENPI 4D NM_002662 NP_002653 3337 SENPI 4D NM_032740 NP_116129 84826 MGC391 4D NM_145266 NP_606369 134492 NUIDCD2 4D NM_006371 NP_006362 10491 CRTAP	solute carrier family 26, mem	SLC26A11	284129	NP_775897	NM_173626		226679_at
4D NM 014554 NP 065369 29843 SENPI 4D NM_002662 NP_002653 5337 PLDI 4D NM_032740 NP_161629 84826 MGCC391 4D NM_145266 NP_660309 134492 NUDCD2	cartilage associated protein	CRTAP	10491	NP_006362	NM_006371		226656_at
4D NM_01454 NP_085369 29843 SENPI 4D NM_002662 NP_082653 5337 PLDI 4D NM_032740 NP_116129 84826 MGCS391	NudC domain containing 2	NUDCD2	134492	NP 660309	NM_145266		226642 s at
4D NM_014554 NP_055369 29843 SENP1 4D NM_002662 NP_002653 5337 PLD1	hypothetical protein MGC539	MGC5391	84826	NP_116129	NM_032740		226639_at
4D NM_014554 NP_055369 29843 SENPI	phospholipase D1, phophatid	PLDI	5337	NP_002653	NM_002662		226636_at
	SUMO1/sentrin specific prot	SENPI	29843	NP_055369	NM 014554		226619 at
4D NM 021075 NP 066553 4731 NDI IEV3	NADH dehydrogenase (ubiquinone) flavoprotein 3, 10	NDUFV3	4731	NP_066553	NM_021075	4D	226616_s_at

similar to RIKEN cDNA 0610008P16 cene	MGC15937	92292	NP 542392	NM 080661	4D	227695 at
zinc finger protein 326	ZNF326	284695	NP_892021	NM 182976	4D	227680_at
Ku70binding protein 3	KUB3	91419	NP_150592	NM_033276	4D	227678_at
zinc finger protein 585A	ZNF585A	199704	NP_689868	NM_152655	4D	227674_at
doublecortin and CaM kinaselike 2	DCAMKL2	166614	NP_689832	NM_152619	40	227666 at
ELAV (embryonic lethal, abnormal vision, Drosophil	ELAVL3	1995	NP_001411	NM_001420	40	227612_at
epithelial stromal interaction 1 (breast)	EPSTII	94240	NP_001002264	NM_001002264	4D	227609_at
KIAA1627 protein	KIAA1627	57721	NP_066012	NM_020961	4D	227601_at
hypothetical protein FLJ12949	FLJ12949	65095	NP_075384	NM_023008	4D	227587 at
hypothetical protein MGC42630	MGC42630	286301	NP_787119	NM_175923	4D	227563_at
PHD finger protein 20like 1	PHF20L1	51105	NP_057102	NM_016018	4D	227523 s_at
Fbox protein 33	FBXO33	254170	NP 976046	NM_203301	4D	227521_at
forkhead box Q1	FOXQ1	94234	NP_150285	NM_033260	45	227475_at
HECT domain and ankyrin repeat containing, E3 ubiq	HACEI	57531	NP_065822	NM_020771	4D	227471_at
hypothetical protein LOC197322	LOC197322	197322	NP_777577	NM_174917	4D	227464 at
EPH receptor A4	EPHA4	2043	NP_004429	NM_004438	4D	227449_at
hypothetical protein FLJ38991	FLJ38991	285521	NP_776188	NM_173827	4D	227442 at
hypothetical protein MGC14595	MGC14595	84294	NP_115710	NM_032334	4D	227402 s at
leucine rich repeat (in FLII) interacting protein	LRRFIP1	9208	NP_004726	NM_004735	4D	227391_x_at
neighbor of COX4	NOC4	10328	NP_006058	NM_006067	4D	227363_s_at
hypothetical protein FLJ90652	FLJ90652	283899	NP_775889	NM_173618	4D	227286_at
neuronal guanine nucleotide exchange factor	NGEF	25791	NP_062824	NM_019850	4D	227240_at
tetraspan 2	TSPAN2	10100	NP_005716	NM_005725	4D	227233_at
KIAA1211 protein	KIAA1211	57482	XP_044178	XM_044178	4D	227230_s_at
ovarian zinc finger protein	HOZFP	152518	NP_694540	NM_152995	4D	227220_at
WNK lysine deficient protein kinase 2	WNK2	65268	NP_006639	NM_006648	4D	227217_at
glutathione Stransferase omega 2	GSTO2	119391	NP_899062	NM_183239	4D	227163_at
chromosome 20 open reading frame 7	C20orf7	79133	NP_077025	NM_024120	4D	227160 s at
hypothetical protein MGC32065	MGC32065	257364	NP_695003	NM_153271	4D	227151_at
BH3 interacting domain death agonist	BID	637	NP_001187	NM_001196	4D	227143 s_at
elongation protein 3 homolog (S. cerevisiae)	ELP3	55140	NP_060561	NM_018091	4D	227075_at
phosphogrycerate kmase i	1001	0630	141 000000	1111 00017	i	- COO

proline argininerich end leucinerich reneat protei	PRELP	5549	NP 002716	NM 002725	40	228224 at
hypothetical LOC389362	LOC389362	389362	XP_374159	XM_374159	4D	228217_s_at
dispatched homolog 1 (Drosophila)	DISPI	84976	NP_116279	NM_032890	4D	228184_at
RPA interacting protein	RIP	84268	NP_115684	NM_032308	4D	228183 s_at
oligodendrocyte transcription factor 1	OLIGI	116448	NP_620450	NM_138983	4D	228170 at
myosin, heavy polypeptide 11, smooth muscle	MYHII	4629	NP_074035	NM_022844	4D	228134 at
hypothetical protein FLJ20280	FLJ20280	54876	NP_060211	NM_017741	4D	228106_at
zinc finger protein 550	ZNF550	162972	NP_694963	NM_153231	4D	228099_at
cAMP responsive element modulator	CREM	1390	NP_001872	NM_001881	4D	228092 at
similar to RIKEN cDNA 1810059G22	LOC374395	374395	NP_955369	NM_199337	4D	228089 x at
calcium channel, voltagedependent, alpha 2/delta s	CACNA2D4	93589	NP_758952	NM_172364	4D	228083_at
hypothetical protein MGC3207	MGC3207	84245	NP_115661	NM_032285	4D	228077 at
Bcell CLL/lymphoma 9like	BCL9L	283149	NP_872363	NM_182557	4D	228065_at
similar to hypothetical protein LOC192734	LOC388886	388886	NP_997527	NM_207644	4D	228064_at
chromosome 9 open reading frame 105	C9orf105	401505	NP_001001790	NM_001001790	4D	228053_s_at
FLJ20758 protein	FLJ20758	55037	NP_060422	NM_017952	4D	228020_at
chromosome 20 open reading frame 58	C20orf58	128414	NP_690603	NM_152864	4D	228017 s at
zinc ribbon domain containing, 1	ZNRDI	30834	NP_055411	NM_014596	4D	228009_x_at
PWWP domain containing 2	PWWP2	170394	NP_612508	NM_138499	4D	227999_at
chromosome 20 open reading frame 149	C20orf149	79144	NP_077275	NM_024299	4D	227994_x_at
postsynaptic protein CRIPT	CRIPT	9419	NP_054890	NM_014171	4D	227942_s_at
exosome component 3	EXOSC3	51010	NP_057126	NM_016042	4D	227916_x_at
CDC42 effector protein (Rho GTPase binding) 5	CDC42EP5	148170	NP_659494	NM_145057	4D	227850_x_at
EPM2A (laforin) interacting protein 1	EPM2AIP1	9852	NP_055620	NM_014805	4D	227847_at
forminlike 3	FMNL3	91010	NP_783863	NM_175736	4D	227844_at
nude nuclear distribution gene E homolog 1 (A. nid	NUEI	24820	NF_060138	NM_01 /088	ŧ	1E_C+0/22
zinc finger protein 605	ZNF605	90462	NP_899061	NM 183238	ŧ	227822 at
zinc finger protein 62 homolog (mouse)	ZFP62	92379	NP_689496	NM_152283	8	227796 at
chromosome 6 open reading frame 157	C6orf157	90025	NP_944602	NM_198920	4D	227790 at
hypothetical protein FLJ90586	FLJ90586	135932	NP_699176	NM_153345	4D	227753 at
C3 and PZPlike, alpha2macroglobulin domain contain	CPAMD8	27151	NP_056507	NM_015692	4D	227721_at
suppressor of cytokine signaling 5	SOCSS	1706	NP_003940	MINI_003933	ŧ	221071 at

collagen (vne XI alpha I	COLITAI	1301	NP_001845	NM 001854	4D	229271 x at
LOC440667	LOC440667	440667	XP_498810	XM_498810	4D	229264 at
lactate dehydrogenase D	LDHD	197257	NP_705690	NM_153486	4D	229241_at
KIAA0251 protein	KIAA0251	23042	NP_055842	NM_015027	4D	229219 s at
solute carrier family 35, member F3	SLC35F3	148641	NP_775779	NM_173508	4D	229065_at
hypothetical protein MGC31967	MGC31967	203260	NP_777583	NM_174923	4D	229063_s_at
nudix (nucleoside diphosphate linked moiety X)type	NODIT.	200035	XP_496393	AM_496393	ŧ	229044_at
zinc finger protein 449	ZNF449	203523	NP 689908	NM_152695	ŧ	228968 at
hypothetical protein FLJ35954	FLJ35954	166968	NP_689835	NM_152622	40	228961_at
S100 calcium binding protein A6 (calcyclin)	S100A6	6277	NP_055439	NM_014624	4D	228923_at
prematurely terminated mRNA decay factorlike	LOC91431	91431	NP_619643	NM_138698	4D	228859_at
hypothetical protein LOC196463	LOC196463	196463	NP_775813	NM_173542	4D	228845 at
solute carrier family 13 (sodiumdependent citrate	SLC13A5	284111	NP_808218	NM_177550	4D	228844_at
KIAA1571 protein	KIAA1571	57683	XP_371590	XM_371590	4D	228749_at
zinc finger, CCHC domain containing 12	ZCCHC12	170261	NP_776159	NM_173798	4D	228715_at
procollagenproline, 2oxoglutarate 4dioxygenase (pr	P4HA3	283208	NP_878907	NM_182904	40	228703_at
chromosome 21 open reading frame 45	C21 orf45	54069	NP_061817	NM_018944	4D	228597_at
FLJ20758 protein	FLJ20758	55037	NP_060422	NM_017952	4D	228590_at
CSRP2 binding protein	CSRP2BP	57325	NP_065397	NM_020536	4D	228544 s_at
CSRP2 binding protein	CSRP2BP	57325	NP_065397	NM_020536	4D	228543_at
sterile alpha motif domain containing 9	SAMD9	54809	NP_060124	NM_017654	4D	228531_at
chromosome 10 open reading frame 86	C10orf86	54780	NP_060085	NM_017615	4D	228506 at
THAP domain containing 8	THAP8	199745	NP_689871	NM_152658	4D	228500_at
6phosphofructo2kinase/fructose2,6biphosphatase	PFKFB4	5210	NP_004558	NM_004567	4D	228499_at
				NM_018675	4D	228393 s at
alphakinase 2	ALPK2	115701	NP_443179	NM_052947	4 D	228367_at
hypothetical protein LOC91942	LOC91942	91942	NP_777549	NM_174889	4D	228355 s at
hypothetical protein MGC16044	MGC16044	91523	NP_612380	NM_138371	4D	228298_at
novel 58.3 KDA protein	LOC91614	91614	NP_631899	NM_139160	4D	228293_at
tudor domain containing 9	TDRD9	122402	NP_694591	NM_153046	4D	228285_at
laminocyclopropanelcarboxylate synthase	PHACS	84680	NP_115981	NM_032592	4D	228264_at

ST8 alphaNacetyIneuraminide	ST8SIA4	7903	NP_005659	NM_005668	4D	230261_at
placentaspecific 9	PLAC9	219348	XP_166090	XM_166090	4D	230246_at
ret finger protein 2	RFP2	10206	NP_005789	NM_005798	4D	230192_at
family with sequence similarity 14, member B	FAM14B	122509	NP 660292	NM_145249	4D	230172_at
similar to Histone H3.3	LOC392533	392533	XP_373368	XM_373368	4D	230144_at
solute carrier family 30 (zinc transporter), membe	SLC30A2	7780	NP_115902	NM_032513	4D	230084_at
BTB (POZ) domain containing 14A	BTBD14A	138151	NP 653254	NM_144653	4D	230050_at
mitochondrial ribosomal protein L43	MRPL43	84545	NP_789763	NM_176793	4D	230026_at
hypothetical protein FLJ21103	FLJ21103	79607	NP_078832	NM_024556	4D	230009_at
tigger transposable element derived 2	TIGD2	166815	NP_663761	NM_145715	4D	229983_at
Ellis van Creveld syndrome 2 (limbin)	EVC2	132884	NP_667338	NM_147127	4D	229974_at
FLJ32363 protein	FLJ32363	375444	NP_940968	NM_198566	4D	229886_at
Kruppellike factor 12	KLF12	11278	NP_009180	NM_007249	4D	229881_at
(ze			1		İ	
nypoinetical EOC145853	VEBH1	79674		AM_096883	5 E	229742 at
similar to RIKEN cDNA 1700027J05 gene	MGC33692	414918	NP_001001/94	NM_001001794	ŧ	229/39 s_at
Tcell activation GTPase activating protein	TAGAP	117289	NP_473455	NM_054114	4D	229723_at
cleavage stimulation factor, 3' preRNA, subunit 3,	CSTF3	1479	NP_001317	NM_001326	4D	229665_at
peroxisomal lon protease	LONP	83752	NP_113678	NM_031490	4D	229663_at
chromosome 18 open reading frame 51	C18orf51	125704	XP_058931	XM_058931	4D	229645_at
iroquois homeobox protein 3	IRX3	79191	NP_077312	NM_024336	4D	229638_at
hypothetical protein FLJ32752	FLJ32752	144132	NP_653267	NM_144666	4D	229631_at
hypothetical protein FLJ39441	FLJ39441	144108	NP_919261	NM_194285	4D	229594_at
Tbox 3 (ulnar manmary syndrome)	ТВХ3	6926	NP_005987	NM_005996	4D	229576_s_at
peroxisomal acylCoA thioesterase 2B	PTE2B	122970	NP_689544	NM_152331	4D	229534_at
fragile X mental retardation, autosomal homolog	FXRI	8087	NP_005078	NM_005087	4D	229519_at
family with sequence similarity 46, member B	FAM46B	115572	NP_443175	NM_052943	4D	229518_at
cytochrome c oxidase subunit Va	COX5A	9377	NP_004246	NM_004255	4D	229426_at
golgi transport 1 homolog A (S. cerevisiae)	GOLTIA	127845	NP_940849	NM_198447	4D	229372_at
immunoglobulin superfamily, member 4C	IGSF4C	199731	NP_660339	NM_145296	4D	229335_at
hypothetical protein MGC2749	MGC2749	79036	NP_076974	NM_024069	4D	229316_at
junctophilin 3	JPH3	57338	NP_065706	NM_020655	4D	229294_at
ribonuclease L (2',5'oligoisoadenylate synthetased	RNASEL	6041	NP 066956	NM_021133	4D	229285_at

chromosome 9 open reading frame 138	C9orf138	158297	NP_714918	NM_153707	4D	231570_at
hypothetical protein MGC35194	MGC35194	93190	NP_689503	NM_152290	4D	231549_at
chromosome 11 open reading frame 1	Cliorfi	64776	NP_073598	NM_022761	4D	231530_s_at
chromosome 20 open reading frame 144	C20orf144	128864	NP_543015	NM_080825	4D	231522_at
gametogenetin	GGN	199720	NP_689870	NM_152657	4D	231420_at
LOC122258	LOC122258	122258	NP_660291	NM_145248	4D	231405_at
carbonic anhydrase XIII	CA13	377677	NP 940986	NM_198584	4D	231270 at
similar to RIKEN cDNA 1110018M03	LOC387758	387758	NP_976249	NM 203371	4D	231001_at
Bcell novel protein 1	BCNP1	199786	NP_775815	NM_173544	4D	230983_at
HCG3 gene	HCG3	414061	NP 001001394	NM_001001394	4D	230953_at
CDC14 cell division cycle 14 homolog B (S. cerevis	CDC14B	8000	NF_201389	NM 03332	ŧ	23066/_at
BCL2associated athanogene 2	BAG2	9532	NP_004273	NM_004282		230879 at
cystatin 9like (mouse)	CST9L	128821	NP_542177	NM_080610		230829_at
SVAP1 protein	IMAGE3451454	116841	NP_444280	NM_053052	4D	230810 at
FLJ37099 protein	FLJ37099	163259	NP 940861	NM_198459	4D	230769_at
hypothetical protein BC014608	LOC128153	128153	NP_620151	NM_138796		230763_at
hypothetical protein MGC33414	MGC33414	257101	NP_775845	NM_173574	4D	230756_at
hypothetical protein LOC146174	LOC146174	146174	NP_775772	NM_173501	4D	230721_at
kinesin family member 9	KIF9	64147	NP_071737	NM_022342	4D	230701_x_at
hypothetical STGC3	STGC3	474171	NP_001006609	NM_001006608	4D	230587_at
hypothetical LOC388480	LOC388480	388480	XP_373779	XM_373779	4D	230574_at
cAMP responsive element modulator	CREM	1390	NP_874391	NM_182722	4D	230511_at
CD5 antigen (p5662)	CD5	921	NP_055022	NM_014207	40	230489_at
similar to RIKEN cDNA 6030419C18 gene	LOC388135	388135	XP_370873	XM_370873	4D	230475_at
hypothetical gene supported by AK124342	LOC400999	400999	XP_379123	XM_379123	4D	230452_at
similar to hypothetical protein 9630041N07	LOC345462	345462	XP_293828	XM_293828	4D	230421_at
inositol polyphosphate5phosphatase F	INPP5F	22876	NP_938145	NM_198331	4D	230363 s at
inositol polyphosphate5phosphatase F	INPP5F	22876	NP_938145	NM_198331	₽Đ	230362_at
transmembrane protein 45B	TMEM45B	120224	NP_620143	NM 138788	4D	230323 s at
chromosome 9 open reading frame 10 opposite strand	C9orf10OS	158293	NP_942138	NM_198841	4D	230308_at
RAB7B, member RAS oncogene family	RAB7B	338382	NP 796377	NM_177403		230266_at
alpha2,8sialyltransfer						

107

piggyBac transposable element derived 3	PGBD3	267004	NP_736609	NM_170753	4D	232287_at
solute carrier family 6, member 15	SLC6A15	55117	NP_060527	NM_018057	4D	232263_at
amyotrophic lateral sclerosis 4	ALS4	23064	NP_055861	NM_015046	4D	232229 at
KIAA1465 protein	KIAA1465	57611	NP_065902	NM_020851	4D	232208_at
palmdelphin	PALMD	54873	NP_060204	NM_017734	4D	232187_at
serine active site containing 1	SERACI	84947	NP_116250	NM_032861	4D	232183_at
hypothetical protein FLJ14129	FLJ14129	79943	NP_112157	NM_030895	4D	232162_at
mediator of RNA polymerase II transcription, subun	MED12L	116931	NP_443728	NM_053002	Đ	232114_at
similar to hypothetical protein Y97E10AL.1	DKFZp761P211	375057	NP_001003665	NM_001003665	4D	232104_at
chromosome X open reading frame 23	CXorf23	256643	NP_938020	NM_198279	4D	232087_at
Nedd4 binding protein 2	N4BP2	55728	NP_060647	NM_018177	4D	231996_at
chromosome 20 open reading frame 160	C20orf160	140706	NP_542192	NM_080625	4D	231991 at
similar to HSPC323	LOC284422	284422	XP_209196	XM_209196	4D	231982_at
PHD finger protein 20like 1	PHF20L1	51105	NP_940915	NM_198513	4D	231967_at
zinc finger protein 529	ZNF529	57711	NP_066002	NM_020951	4D	231940 at
densityregulated protein	DENR	8562	NP_003668	NM_003677	4D	231896 s_at
RNA (guanine9) methyltransferase domain containing	RG9MTD2	93587	NP_689505	NM_152292	4D	231877_at
kinesin family member 21A	KIF21A	55605	NP_060111	NM_017641	4D	231875_at
inhibitor of growth family, member 3	ING3	54556	NP_938008	NM_198267	4D	231863_at
tubulin, delta 1	TUBDI	51174	NP_057345	NM_016261	4D	231853_at
alanyltRNA synthetase like	AARSL	57505	NP_065796	NM_020745	4D	231845_at
KIAA1272 protein	KIAA1272	57186	XP_046600	XM_046600	4D	231826_at
FLJ10378 protein	FLJ10378	55132	NP_060548	NM_018078	4D	231824_at
noggin	NOG	9241	NP_005441	NM_005450	4D	231798_at
distalless homeo box 3	DLX3	1747	NP_005211	NM_005220	4D	231778_at
member				1		
tumor necrosis factor receptor superfamily.	TNFRSF10A	8797	NP_003835	NM_003844	4D	231775_at
upstream transcription factor 1	USFI	7391	NP_009053	NM_007122	4D	231768_at
zona pellucida głycoprotein 4	ZP4	57829	NP_067009	NM_021186	4D	231756 at
winglesstype MMTV integration site family, member	W IN IS	/4/3	NF_110360	NINI_030733	ŧ	10 C+/ 1 CT
tissue inhibitor of metalloproteinase 2	IIMP2	7077	NP 003240	NM 030353	ŧ	231742 s at
guanylate binding protein 1, interferoninducible,	GBPI	2633	NP_002044	NM_002053	6	231578 at

CXYorf1related protein	FLJ25222	374666	NP_954632	NM_199163	4D	233929 x at
SLITROBO Rho GTPase activating protein 1	SRGAPI	57522	NP_065813	NM_020762	4D	233888 s at
chromosome 20 open reading frame 43	C20orf43	51507	NP_057491	NM_016407	4D	233842 x at
ATbinding transcription factor 1	ATBF1	463	NP_008816	NM_006885	4D	233751_at
keratin associated protein 94	KRTAP94	85280	NP_149461	NM_033191	4D	233640 x at
signalinduced proliferationassociated 1 like 2	SIPAIL2	57568	NP_065859	NM 020808	4D	233587 s at
chromosome 20 open reading frame 149	C20orf149	79144	NP_077275	NM_024299	4D	233571_x_at
cofilin 2 (muscle)	CFL2	1073	NP_068733	NM_021914	4D	233496_s_at
hypothetical protein LOC96597	LOC96597	96597	XP_378655	XM_378655	4D	233483_at
zinc finger protein 226	ZNF226	7769	NP 057528	NM_016444	4D	233461_x_at
polymerase (DNA directed) nu	POLN	353497	NP_861524	NM_181808	4D	233454_at
DEP domain containing 2	DEPDC2	80243	NP_079146	NM_024870	4D	233408_at
	PCNT2	5116	NP_006022	NM_006031	4D	233387 s_at
polymerase (RNA) I polypeptide B, 128kDa	HINTOP	84172	NP_061887	NM_019014	4D	233341 s_at
churchill domain containing 1	CHURC1	91612	NP_660148	NM_145165	4D	233268 s_at
hypothetical protein FLJ23322	FLJ23322	80020	NP_079231	NM_024955	4D	233250_x_at
general transcription factor IIIC, polypeptide 5,	GTF3C5	9328	NP_036219	NM_012087	4D	233173_x_at
UDPNacetylalphaDgalactosamine:polypeptide Nacetylg	GALNTL5	168391	NP_660335	NM_145292	4D	233150_at
chromosome 20 open reading frame 142	C20orf142	128486	XP_371399	XM_371399	4D	233061_at
chromosome X open reading frame 44	CXorf44	90736	NP_612371	NM_138362	4D	232990_at
hypothetical protein FLJ20444	FLJ20444	403323	NP_982274	NM_203449	4D	232897_at
chromosome 20 open reading frame 166	C20orf166	128826	NP_848558	NM_178463	4D	232892_at
dedicator of cytokinesis 8	DOCK8	81704	NP_982272	NM_203447	4D	232843 s_at
PWWP domain containing 1	PWWPI	154150	NP_612641	NM_138574	4D	232680_at
hypothetical gene supported by AK022396; AK097927	LOC400946	400946	XP_379079	XM_379079	4D	232664_at
APG16 autophagy 16like (S. cerevisiae)	APG16L	55054	NP_060444	NM_017974	4D	232612_s_at
WAP fourdisulfide core domain 3	WFDC3	140686	NP_542181	NM_080614	4D	232602_at
hypothetical protein MGC27466	MGC27466	127396	NP_689586	NM_152373	4D	232563_at
leucine rich repeat and fibronectin type III domai	LRFNI	57622	XP_290842	XM_290842	4D	232486_at
chromosome 10 open reading frame 71	C10orf71	118461	NP_955629	NM_199459	4D	232456_at
hypothetical protein FLJ21439	FLJ21439	80208	NP_079413	NM_025137	4D	232403_at
PDZ and LIM domain 2 (mystique)	PDLIM2	64236	NP_789847	NM_176871	4D	232302_at

NYI-4308244v1

TAF3 RNA polymerase II, TATA box binding	TAF3	83860	XP_291729	XM_291729	4D	235119_at
hook homolog 3 (Drosophila)	ноокз	84376	NP_115786	NM_032410	40	235114 x at
HRASlike suppressor 3	HRASLS3	11145	NP 009000	NM_007069	4	235110_at
Fbox and leucinerich repeat protein 20	FBXL20	84961	NP_116264	NM_032875	4D	235089_at
nuclear domain 10 protein	NDP52	10241	NP_005822	NM_005831	4D	235076_at
transmembrane protein 41A	TMEM41A	90407	NP_542383	NM_080652	4D	235037_at
similar to Zinc finger protein 58 (Zfp58) (Zinc ti	LOC402524	402524	XP_379850	XM_379850	4D	235027_at
TAF4b RNA polymerase II, TATA box binding protein	TAF4B	6875	XP_290809	XM_290809	4	235020_at
BardetBiedl syndrome 7	BBS7	55212	NP_789794	NM_176824	4D	235007_at
valosincontaining protein (p97)/p47 complexinterac	VCIP135	80124	NP_079330	NM_025054	40	234988_at
hypothetical protein LOC143458	LOC143458	143458	NP_777562	NM_174902	4D	234985 at
similar to Zincalpha2glycoprotein precursor (Znalp	LOC401393	401393	XP_376658	XM_376658	#	234955_at
deoxynucleotidyltransferase, terminal, interacting	DNTTIPI	116092	NP_443183	NM_052951	4D	234942 s_at
ribosomal protein L7a	RPL7A	6130	NP_000963	NM_000972	\$	234873 x at
transient receptor potential cation channel, subfa	TRPM6	140803	NP_060132	NM_017662	4D	234864_s_at
G2 protein	G2	25758	XP_039515	XM_039515	Ð	234784 at
KIAA1596	KIAA1596	57697	XP_048128	XM_048128	4D	234733 s_at
poly (ADPribose) polymerase family, member 6	PARP6	56965	NP 064598	NM_020213	4D	234710_s_at
similar to bA182L21.1 (novel protein similar to hy	LOC399788	399788	XP_374817	XM_374817	4D	234695_x_at
olfactory receptor, family 52, subfamily D, member	OR52D1	390066	NP_001005163	NM_001005163	ŧ	234617_at
likely ortholog of mouse NOBOX	NOBOX	135935	XP_069612	XM_069612	4D	234519 at
sperm associated antigen 4like	SPAG4L	140732	NP_542406	NM_080675	4D	234504_at
cytotoxic Tlymphocyteassociated protein 4	CTLA4	1493	NP_005205	NM_005214	4b	234362_s_at
zinc finger protein 326	ZNF326	284695	NP_892020	NM_182975	4D	234093 at
serine/threonine kinase 36 (fused homolog, Drosoph	STK36	27148	NP_056505	NM_015690	40	234005_x_at
pleckstrin homology domain containing, family G	PLEKHG2	64857	NP_073746	NM_022835	4D	233986_s_at
espin	ESPN	83715	NP_113663	NM_031475	4D	233979 s at
CGI09 protein	CG109	51605	NP_057023	NM_015939	4D	233970 s at
cystatin 11	CST11	140880	NP_570612	NM_130794	1	233968 at

NY14308244v1

IQ motif containing F2	IQCF2	389123	NP_982248	NM_203424	4D	236767_at
hypothetical protein FLJ39485	FLJ39485	285603	NP_787116	NM_175920	4D	236728 at
x 009 protein	MDS009	56986	NP_064619	NM_020234	4D	236649_at
ribosomal protein S27 (metallopanstimulin 1)	RPS27	6232	NP_001021	NM_001030	4D	236621_at
docking protein 6	DOK6	220164	NP_689934	NM_152721	4D	236290_at
eukaryotic translation initiation factor 3, subuni	EIF3S9	8662	NP_003742	NM_003751	4D	236274 at
zinc finger protein 346	ZNF346	23567	NP_036411	NM_012279	4D	236267_at
multimerin 2	MMRN2	79812	NP_079032	NM 024756	4D	236262_at
phosphatase, orphan 1	PHOSPHOI	162466	NP 848595	NM_178500	4D	236218_at
KIAA1853 protein	KIAA1853	84530	NP_919262	NM_194286	4D	236151_at
actin binding LIM protein family, member 2	ABLIM2	84448	NP_115808	NM_032432	4D	236087_at
X antigen family, member 3	XAGE3	170626	NP_573440	NM_133179	4D	236040 at
chromosome 10 open reading frame 78	C10orf78	119392	NP_001002759	NM_001002759	4D	236027 at
vesicle transport through interaction with tSNAREs	VTIIA	143187	NP_660207	NM_145206	4D	235923_at
guanylate binding protein 4	GBP4	115361	NP_443173	NM_052941	4D	235574_at
hypothetical protein MGC40214	MGC40214	137682	NP_689629	NM_152416	4D	235509_at
ankyrin repeat and SOCS boxcontaining 5	ASB5	140458	NP_543150	NM_080874	4D	235503_at
fucosyltransferase 10 (alpha (1,3) fucosyltransfer	FUT10	84750	NP_116053	NM_032664	4D	235472_at
TruB pseudouridine (psi) synthase homolog 1 (E. co	TRUBI	142940	NP_631908	NM_139169	4D	235447_at
shugoshinlike 2 (S. pombe)	SGOL2	151246	NP_689737	NM_152524	4D	235425_at
similar to RP2 protein, testosteroneregulated	LOC390916	390916	XP_372723	XM_372723	4D	235384_at
ELLP3030	UNQ3030	375387	NP_940967	NM_198565	4D	235359_at
UBX domain containing 4	UBXD4	165324	NP_859064	NM_181713	4D	235327 x at
RNA (guanine9) methyltransferase domain containing	RG9MTD3	158234	NP_659401	NM_144964	4D	235303_at
eukaryotic translation initiation factor 5A2	EIF5A2	56648	NP_065123	NM_020390	40	235296_at
bone morphogenetic protein 8b (osteogenic protein	ВМР8В	656	NP_001711	NM_001720	4D	235275_at
DCP2 decapping enzyme homolog (S. cerevisiae)	DCP2	167227	NP_689837	NM_152624	4D	235258_at
folliculin	FLCN	201163	NP_653207	NM_144606	4D	235250_at
hypothetical protein FLJ90709	FLJ90709	153129	NP_775785	NM_173514	4D	235241_at
similar to hepatocellular carcinomaassociated anti	LOC151194	151194	NP_660323	NM_145280	4D	235177_at
protein (

hypothetical protein FLJ20582	FLJ20582	54989	NP_054825	NM_014106	4D	238687 x at
WD repeat domain 36	WDR36	134430	NP_644810	NM_139281	4D	238677_at
chromosome 14 open reading frame 150	C14orf150	112840	NP_542397	NM_080666	4D	238650 x at
hypothetical protein FLJ21657	FLJ21657	64417	NP_071928	NM_022483	4D	238635_at
hypothetical protein MGC2474	MGC2474	65988	NP_076420	NM_023931	4D	238606_at
Cblinteracting protein Sts1	STS1	84959	NP_116262	NM_032873	4D	238587 at
hypothetical protein MGC39389	FLJ32384	124790	NP_653209	NM_144608	4D	238547_at
synleurin	SLRN	353281	NP_852607	NM_181506	4D	238488_at
similar to Pl3kinaserclated kinase SMG1 isoform 1;	LOC440345	440345	XP_496125	XM_496125	4D	238449_at
citrate lyase beta like	CLYBL	171425	NP_996531	NM_206808	4D	238440_at
carbonic anhydrase VBlike	CASBL	340591	XP_291346	XM_291346	4D	238435_at
hypothetical protein FLJ33069	FLJ33069	137835	NP_653250	NM_144649	4D	238429_at
hypothetical LOC339541	MGC33556	339541	NP_001004307	NM_001004307	4D	238365_s_at
nuclear receptor coactivator 6 interacting protein	NCOA6IP	96764	NP_079107	NM_024831	4D	238346_s_at
TEA domain family member 2	TEAD2	8463	NP_003589	NM_003598	4D	238323_at
TEA domain family member 2	TEAD2	8463	NP_003589	NM_003598	4D	238322_s_at
FB14 protein	FB14	404201	NP_995329	NM_205857	4D	238081_at
retinol binding protein 7, cellular	RBP7	116362	NP_443192	NM_052960	4D	238066_at
hypothetical protein FLJ32028	FLJ32028	201799	NP_689893	NM_152680	4D	238063_at
hypothetical LOC400849	LOC400849	400849	XP_378988	XM_378988	4D	237806_s_at
chromosome 14 open reading frame 68	C14orf68	283600	NP_997000	NM_207117	4D	237765_at
hypothetical LOC197387	LOC197387	197387	XP_117030	XM_117030	4D	237730_at
hypothetical protein FLJ10569	FLJ10569	55174	NP_060612	NM_018142	4D	237504_at
UDPGal:betaGal beta 1,3galactosyltransferase polyp	B3GALT7	374907	NP_940942	NM_198540	4D	237338_at
transglutaminase 7	TGM7	116179	NP_443187	NM_052955	4D	237288_at
protein kinase, interferoninducible double strande	PRKRA	8575	NP_003681	NM_003690	4D	237107_at
chromosome 20 open reading frame 70	C20orf70	140683	NP_542141	NM_080574	4D	237099_at
CWF19like 2, cell cycle control (S. pombe)	CWF19L2	143884	NP 689647	NM_152434	4D	237040_at
chromosome 6 open reading frame 128	C6orf128	221468	NP_660359	NM_145316	4D	237016_at
hypothetical protein MGC20470	MGC20470	143630	NP_659490	NM_145053	4D	236965_at
hypothetical protein MGC50372	MGC50372	253143	NP_775837	NM_173566	4D	236941_at
testis expressed sequence 13A	TEX13A	56157	NP_112564	NM_031274	4D	236848_s_at

238722_x_at	4D	NM_198990	NP_945341	222236	NAPEPLD	Nacylphosphatidylethanolaminehydrolyzing phospholi
238760_at	4D	NM_003680	NP_003671	8565	YARS	tyrosyltRNA synthetase
238817_at	4D	NM_015347	NP_056162	23504	KIAA0318	RIM binding protein 2
238858_at	40	NM_052864	NP_443096	92610	TIFA	TRAFinteracting protein with a forkheadassociated
238940_at	4D	NM 007249	NP_009180	11278	KLF12	Kruppellike factor 12
238974_at	4D	NM_153689	NP_710156	205327	FLJ38973	hypothetical protein FLJ38973
238982 at	4D	NM_003677	NP_003668	8562	DENR	densityregulated protein
238996_x_at	4D	NM_000034	NP_000025	226	ALDOA	aldolase A, fructosebisphosphate
239042_at	4D	NM_018128	NP_060598	55720	FLJ10534	hypothetical protein FLJ10534
239081_at	4D	XM_498525	XP_498525	440064	LOC440064	LOC440064
239186_at	4D	XM_376463	XP_376463	221756	MGC39372	hypothetical protein MGC39372
239196_at	4D	NM_144590	NP_653191	118932	ANKRD22	ankyrin repeat domain 22
239205_s_at	4D	NM_000573	NP_000564	1378	CRI	complement component (3b/4b) receptor 1, including
239212 at	₽ Đ	NM_032860	NP_116249	84946	C6orf93	chromosome 6 open reading frame 93
239261_s_at	4D	NM_006587	NP_006578	10699	CORIN	corin, serine protease
239335_at	4D	NM_198526	NP_940928	374655	DKFZp547K111 3	hypothetical protein DKFZp547K1113
239377_at	4D	NM_032325	NP_115701	84285	MGC11102	hypothetical protein MGC11102
239382_at	4D	NM_000195	NP_000186	3257	ISAH	HermanskyPudlak syndrome 1
239430_at	4D	NM_198541	NP_940943	374918	IGFLI	insulin growth factorlike family member 1
239462_at	4D	XM_496317	XP_496317	342909	4	zinc finger protein 284
239481_at	4D	NM_173698	NP_775969	286499	FLJ37659	hypothetical protein FLJ37659
239598 s_at	4D	NM_017839	NP_060309	54947	FLJ20481	hypothetical protein FLJ20481
239624_at	4D	XM_498906	XP_498906	440885	LOC440885	LOC440885
239635_at	4D	NM_032886	NP_116275	84972	MGC15912	hypothetical protein MGC15912
239648 at	4D	NM_173475	NP_775746	123879	MGC48972	hypothetical protein MGC48972
239738_at	4D	NM_053281	NP_444511	117154	DACH2	dachshund homolog 2 (Drosophila)
240027_at	4D	NM_004664	NP_004655	8825	LIN7A	lin7 homolog A (C. elegans)
240082 s at	4D	NM_175734	NP_783861	201243	LOC201243	hypothetical protein LOC201243
240159_at	4D	NM_021082	NP_066568	6565	SLC15A2	solute carrier family 15 (H+/peptide transporter),
240359_at	4D	NM_178466	NP_848561	128861	C20orf71	chromosome 20 open reading frame 71
240603 s at	4D	NM_080663	NP_542394	112479	MGC16943	similar to RIKEN cDNA 4933424N09 gene

DKFZP4341116 protein	DKFZP4341116	25962	NP_056311	NM_015496	4D	243927 x at
rab3 G1 Paseactivating protein, noncatalytic subuni	RABJUAPIO	79/07	NF_030340	NE 012414	ŧ	240071_at
collagen and calcium binding EGF domains	CCBEI	14/3/2	NP 03/210	NM_133439	ŧ	243805 at
MARROW ZI	MOCIZOIS	01000	NF_271020		đ	213001_40
solute carrier family 1 (glutamate transporter), m	SLCIA/	2100	799900 AN) ŧ	243625 at
REVIlike (yeast)	REVIL	51455	NP 05/400	NM 016516	ŧ	243592 at
APG10 autophagy 10tike (S. cercvisiae)	APG10L	83734	NP_113670	NM 031482	ŧ	2434//_at
CasBrM (murine) ecotropic retroviral transforming	CBL	807	NF_005179	NIV. 003188	ŧ	240470_at
zinc finger protein 214	ZNF214	7761	NP_03/381	NM 013249	ŧ	243437 s at
zinc finger protein 214	ZNF214	7761	NP_037381	NM_013249	4	243456 at
hypothetical protein LOC339290	LOC339290	339290	XP_375527	XM_375527	4D	243426_at
mabal	KIAA1324	57535	NP_065826	NM_020775	4D	243349_at
KIBRA protein	KIBRA	23286	NP_056053	NM_015238	4D	243290_at
FLJ35093 protein	FLJ35093	374986	NP_940951	NM_198549	4D	243042_at
ADPribosylation factorlike 8	ARL8	221079	NP_848930	NM_178815	4D	242727_at
hypothetical protein FLJ13305	FLJ13305	84140	XP_291019	XM_291019	4D	242584_at
TNF receptorassociated factor 4	TRAF4	9618	NP_004286	NM_004295	4D	242473_at
hypothetical protein LOC199899	LOC199899	199899	XP_378866	XM_378866	4D	242284 at
poly(A) polymerase beta (testis specific)	PAPOLB	56903	NP_064529	NM_020144	4D	242158_at
orthodenticle homolog 2 (Drosophila)	OTX2	5015	NP_068374	NM_021728	4D	242128_at
chromosome 6 open reading frame 152	C6orf152	167691	NP_859065	NM_181714	4D	242006_at
checkpoint suppressor I	CHESI	1112	NP_005188	NM_005197	4D	241984_at
WD repeat domain 4	WDR4	10785	NP_387510	NM_033661	4D	241937 s at
calmodulin binding transcription activator 1	CAMTAI	23261	NP_056030	NM_015215	4Đ	241882_at
sodium channel, nonvoltagegated 1, gamma	SCNNIG	6340	NP_001030	NM_001039	4D	241436 at
hypothetical protein FLJ34443	FLJ34443	285464	NP_787114	NM_175918	4D	241408 at
similar to hypothetical protein	LOC387816	387816	XP_370664	XM_370664	4D	241360 at
hypothetical LOC283034	LOC283034	283034	XP_210860	XM_210860	4D	241337_at
forkhead box N4	FOXN4	121643	NP_998761	NM_213596	4D	241009_at
keratin associated protein 193	KRTAP193	337970	NP_853640	NM_181609	4D	240967_at
fibroblast growth factor receptor 2 (bacteriaexpre	FGFR2	2263	NP_075262	NM 022973	4D	240913 at

DEAD (AsnGlijAlaAsn) hov polypentide 28	DDX28	55794	NP 060850	NM 018380	4D	40255 at
SH2B homolog	SH2B	25970	NP_056318	NM_015503	4D	40149_at
interleukin 1, beta	IL1B	3553	NP_000567	NM_000576	4D	39402_at
extra spindle poles like 1 (S. cerevisiae)	ESPL1	9700	NP_036423	NM_012291	4D	38158_at
parvin, beta	PARVB	29780	NP_001003828	NM_001003828	4Đ	37966 at
prolyl endopeptidase	PREP	5550	NP_002717	NM_002726	4D	37950_at
surfactant, pulmonaryassociated protein B	SFTPB	6439	NP_000533	NM_000542	4D	37004_at
vmaf musculoaponeurotic fibrosarcoma oncogene homo	MAFF	23764	NP_036455	NM_012323	ŧ	56/11_at
Nsulfoglucosamine sulfohydrolase (sulfamidase)	SGSH	6448	NP_000190	NM_000199	4D	35626_at
FLN29 gene product	FLN29	10906	NP_00691	NM_006700	4D	35254_at
reticulon 2	RTN2	6253	NP_005610	NM_005619	4D	34408_at
peroxisomal biogenesis factor 14	PEX14	5195	NP_004556	NM_004565	4D	33760_at
interferon stimulated gene 20kDa	ISG20	3669	NP_002192	NM_002201	4D	33304_at
myosin IC	MYOIC	4641	NP_203693	NM_033375	4D	32811_at
Nedd4 binding protein 1	N4BPI	9683	NP_694574	NM_153029	4D	32069_at
hypothetical protein BC002942	BC002942	91289	NP_149977	NM_033200	4D	31837_at
DEAD (AspGluAlaAsp) box polypeptide 49	DDX49	54555	NP_061943	NM_019070	45	31807_at
bromodomain and WD repeat domain containing	BRWD3	254065	NP_694984	NM_153252	4D	244738 at
nuclear transcription factor Y, beta	NFYB	4801	NP_006157	NM_006166	4D	244704_at
hypothetical protein FLJ36674	FLJ36674	284040	NP_775893	NM_173622	4D	244698_at
cytochrome c, somatic	CYCS	54205	NP_061820	NM_018947	4D	244546_at
hypothetical LOC200624	LOC200624	200624	XP_117257	XM_117257	4D	244518_at
hypothetical protein MGC27466	MGC27466	127396	NP_689586	NM_152373	4D	244398_x_at
hypothetical protein FLJ31340	FLJ31340	222223	NP_689961	NM_152748	4D	244317_at
THAP domain containing 5	THAP5	168451	NP_872335	NM_182529	4D	244190_at
muskelin 1, intracellular mediator containing kelc	MKLNI	4289	NP_037387	NM_013255	4D	244171_at
gammaaminobutyric acid (GABA) A receptor, alpha 1	GABRA1	2554	NP_000797	NM_000806	D	244118_at
hypothetical protein FLJ35382	FLJ35382	163859	NP_689821	NM_152608	4D	244103_at
protein phosphatase 1K (PP2C domain containing)	PPMIK	152926	NP_689755	NM_152542	4D	244011_at
ATPbinding cassette, subfamily B (MDR/TAP), member	ABCB1	5243	NP_000918	NM_000927	Đ	243951_at
r Amort of incident	INAGI	144	141 0/5550	141AC 0720-14	-	27.770 at

					37)
40288 at 4D	NM_020676	NP_065727	57406	ABHD6	abhydrolase domain containing 6
47550 at 4D	NM_021020	NP_066300	11178	LZTSI	leucine zipper, putative tumor suppressor
48030 i at 4D	NM_032385	NP_115761	10826	C5orf4	chromosome 5 open reading frame 4
49077_at 4D	NM_016147	NP_057231	51400	PMEI	protein phosphatase methylesterase!
51228_at 4D	NM_203390	NP 976324	389677	LOC389677	similar to RIKEN cDNA 3000004N20
53968_at 4D	NM_030628	NP_085131	80789	KIAA1698	KIAA1698 protein
53987_at 4D	NM_020850	NP_065901	57610	RANBPI0	RAN binding protein 10
54037_at 4D	NM_022081	NP_071364	89781	HPS4	HermanskyPudlak syndrome 4
55662_at 4D	NM_024541	NP_078817	79591	C10orf76	chromosome 10 open reading frame 76
55872_at 4D	NM_020713	NP_065764	57473	KIAA1196	KIAA1196 protein
57516_at 4D	NM_033410	NP_219363	92595	MGC13138	hypothetical protein MGC13138
57532_at 4D	NM_004422	NP_004413	1856	DVL2	dishevelled, dsh homolog 2 (Drosophila)
57588_at 4D	NM_020689	NP_065740	57419	SLC24A3	solute carrier family 24
					(sodium/potassium/calcium
	NM_018130	NP_060600	55164	SHQI	SHQ1 homolog (S. cerevisiae)
64432 at 4D	NM_016534	NP_057618	51275	FLJ39616	apoptosisrelated protein PNASI
65493 at 4D	NM_022070	NP_071353	63897	ABCI	amplified in breast cancer 1
74694_s_at 4D	NM_024816	NP_079092	79874	RABEP2	rabaptin, RAB GTPase binding effector protein 2
AFFX- NM_007315	15 NP_009330		6772	STATI	signal transducer and activator of transcription 1
97934D					
AFFX- NM_007315 HUMISGF3A/M 97934D	NP_009330		6772	STATI	signal transducer and activator of transcription 1